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                                                                                                                                                                Use of thioredoxin for treatment of ma
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12-JUN-1991
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                                                                                                                                                                co-factor(s) e.g. interferon(s) etc
e.g. B lymphocytic leukaemia, colon
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Abb11846
Add46716
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Ade55400
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Add
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Aag22068
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add missing
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RESULT 2
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                                         CA AAW09052-W09054 are motifs that are present in thioredoxin or enzymes CC containing thioredoxin domains, e.g. protein-disulphide oxidoxeductases. CC These enzymes are used in a method for producing biologically active, CC correctly branched, disulphide-bonded proteins. The method involves CC reacting the corresponding non-branched, reduced protein with a protein-cC disulphide oxidoreductase enzyme that allows correct disulphide bonding, CC in the presence of glutaredoxin and redox buffer (glutathione disulphide) CC with a redox potential of -270 to -140 mV, to give protein with desired CC biological activity. The method is used for producing human proteins in CC bacteria (e.g. Escherichia coli) in the correct conformation. Proteins CC congulation factors, interleukins, pancreatic mammalian trypsin inhibitors, interferons, remnin, CC prolactin and human alpha-1 trypsin inhibitors can be made in prokaryotic cells in a biologically active form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 4
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                                                                                                                                                                                                                                                                                                                      Preparing correctly branched proteins contg. di:sulphide linkages comprises reacting non-branched reduced protein with protein di:su oxido:reductase enzyme in glutaredoxin and redox buffer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thioredoxin; protein folding; branched; disulphide bond; prokaryotic recombinant production; biologically active; protein disulphide oxidoreductase; glutaredoxin; redox buffer
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                                                                                                                                                                                                                                                                                                                                                                                                       Ljung
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HOLM/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW09052;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW09052 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  field.)
                                        Sequence
                                                                                                                                                                                                                                                                                              Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                    (ASLU/)
                                                                                                                                                                                                                                                                                            2; Page 16; 22pp; Swedish.
                                                                                                                                                                                                                                                                                                                                                                                                        J, Holmgren
                                                                                                                                                                                                                                                                                                                                                                                                                                  ) LJUNG J.
) HOLMGREN A.
) ASLUND F.
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100.0%; Score 31; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                          Aslund
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Pred. No.
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                                                                                                                                                                                                                                                                                                    The present polypeptide is a conserved motif found in the Trx-homology domain of the thioredoxin (Trx) family of proteins. The motif is cimportant for catalytic activity. The N-terminal region of PKC-interacting cousin of thioredoxin (PICOT) shows homology to the Trx-homology domain but it lacks the conserved motif and contains instead an Ala-Pro-Gln-Cys motif. PICOT modulates the function of PKCtheta and inhibits activation of c-Jun N-terminal kinase (JNK) and the transcription factors AP-1 and MF-kappaB. These polypeptides are involved in the TCR/CD28 signaling cascade which leads to IL-2 production. PICOT also modulates activity of the Trx system. PICOT may be useful for ameliorating physiological conditions, e.g. an inflammation or stress response, associated with these polypeptides. The cDNA encoding PICOT solated using yeast two-hybrid screening. PKCtheta polypeptides were screened for binding to polypeptides encoded by a Jurkat T lymphoma cDNA ibrary. The two longest cDNAs from the PKCtheta-interacting positive clones were sequenced and the putative 37.5 kDa polypeptide encoded by
                                                                                                                      Query Match
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Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A PKC-interacting cousin of Trx (PICOT) polypeptides and the nucleic acids that encode it, useful e.g. for modulating Jun N-terminal kinase, AP-1 and/or nuclear factor-kappa-B expression and activity.
                                                                                                                                                                                                                                                      Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-442379/38.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200036083-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human Trx-homology domain conserved motif
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 43; 68pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (LJOL-) LA JOLLA INST ALLERGY & IMMUNOLOGY.
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                                                       CGPC 5
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                                                                                                                         Conservative
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                                                                                                                                                100.0%;
                                                                                                                      Score 31; DB 3;
Pred. No. 2e+06;
; Mismatches
                                                                                                                                                                                    DB 3;
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                                                                                                                                                                                    Length 4;
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RESULT 4

05-AUG-1998

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RESULT 5
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ID ABP5
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Matches 4
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                                 CN1189539-A
                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                 Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Col 15; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Saccharomyces; mammalian; fusion protein; interactor peptide; conformation-constraining protein; DNA binding moiety; thioredgene activating moiety; protein interaction; gene purification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY49348 standard;
                                                                               Thioredoxin;
                                                                                                      Thioredoxin related peptide #2.
                                                                                                                                                     ABP56403;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Population of Saccharomyces and/or mammalian cells comprising recombinant DNA encoding fusion proteins, useful for detecting protein interactions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Escherichia coli.
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                                                                                                                                                                           ABP56403 standard;
                                                                                                                                                                                                                                                                                                                                                       polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUL-1994;
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ilarity 100.0%;
Conservative (
                                                                               TrxA; fusion protein; allogenic protein.
                                                                                                                              (first entry)
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                                                                                                                                                                          peptide;
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                                                                                                                                                                                                                                                                                                        Length 4;
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RRESULT 6
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XX AA022
XX Redc
XX Anti
KW Anti
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KW Heli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a DNA molecule of a fusion protein which is composed of the DNA sequence similar to encoded thioredoxin in structure and the DNA sequence of an encoded allogenic protein. The soluble expression of the fusion protein is efficient. The expressed fusion protein can be easily purified as several amino acid sequences are inserted at active sites of the thioredoxin gene sequence, and hydrolysed by a specific protease as the junction between the two encoded DNA sequences can be cut by the specific protease, obtaining a recombined allogenic protein the same as the natural protein in one-class structure. The present sequence represents a peptide which is given in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                               chronic disease; autoimmune disease; asthma; septic shock; lung fibrosis glomerulonephritis; atherosclerosis; autoimmune encephalomyelitis; soft tissue injury; reduction catalyst; chronic inflammatory disease; systemic inflammatory response; NF-kappaB-mediated inflammatory disease; inducible surface expression; CD44; intercellular adhesion molecule; ICAM-1; NF-kappaB activation; redox active peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antiinflammatory; immunosuppressive; antiasthmatic; antibacterial; nephrotropic; antiarteriosclerotic; antiheumatic; antiarthritic; Helicobacter pylori protein; inhibitor; activator; NF-kappaB; rheumatoid; inflammation; inflammatory bowel disease; autoimmune arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-JAN-1997;
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   WPI; 2002-627466/67
                                                         Windle HJ,
                                                                                                                                                                           30-JAN-2001;
                                                                                                                                                                                                                                 30-JAN-2002; 2002WO-IE000011
                                                                                                                                                                                                                                                                                          08-AUG-2002
                                                                                                                                                                                                                                                                                                                                                   WO200260933-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Redox active peptide sequence CGPC
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                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BIOL-)
                                                                                                                  (QUEE-) QUEEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 capc s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 2 (Disclosure); 14pp;
                                                         Dermot
                                                                                                                                                                           2001IE-00000069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                     ELIZABETH COLLEGE DUBLIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97CN-00100362
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                                                         <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
                                                         Dermot 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 31; DE
Pred. No. 2e+
0; Mismatches
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                                                         Abdel-Latif MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fibrosis;
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ABG32509

ABG32509

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AC ABG3

XX

AC ABG3

XX

M. 15-1

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M. 10-1

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MO2

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MO3

PFF 07-

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MO4

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a Helicobacter pylori protein, its derivatives, cc fragments, mutants and variants capable of inhibiting the activation of confidence of fragments, mutants and variants capable of inhibiting the activation of confidence of the amino acids defined in the specification. The protein is useful for preventing and/or treating confidence of inflammatory bowel disease, rheumatoid/autoimmune confidence of inflammatory of inflammatory autoimmune encephalomyelitis, and confidence of inflammatory of inflammatory confidence of inflammatory confidence of inflammatory confidence of inflammatory response syndrome or other confidence of particularly systemic inflammatory response syndrome or other confidence of inflammatory disease states, down regulates the confidence of inflammatory response syndrome or other confidence of inflammatory disease states, down regulates the confidence of inflammatory response syndrome or other confidence of inflammatory response syndrome or other confidence of inflammatory response syndrome or other confidence of inflammatory disease states, down regulates the confidence of inflammatory response syndrome or other confidence inflammatory response syndrome or other confidence inflammatory response syndrome or other confidence inflammatory response regulates the confidence of inflammatory response represents of inflammatory response response response response res
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Best Local S
Matches 4
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                                                                                                                                                                                                                                                   07-FEB-2001; 2001US-0267328P.
18-SEP-2001; 2001US-0322989P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               M. tuberculosis thioredoxin active site peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                               07-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                   15-AUG-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mycobacterium tuberculosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 attenuated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tuberculostatic; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABG32509;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABG32509 standard; peptide;
                                                                           WPI; 2002-643387/69
                                                                                                                                                                                                                                                                                                                                                                                                                               WO200262298-A2
                                                                                                                                                                             (USGO )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Page 22; 34pp; English.
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                                                                                                                                                                                                       UNIV VANDERBILT
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                                                                                                                            DS,
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                                                                                                                                                                             DEPT VETERANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 intracellular bacterium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TRX; tuberculosis; antibacterial; active site;
                                                                                                                                                                                                                                                                                                                                 2002WO-US003451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                            Bochan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         superoxide dismutase; SOD; bacterial infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 31; DB
Pred. No. 2e+
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Modifying a bacterium to enhance immunogenicity, as vaccines preventing bacterial infections, e.g. tuberculosis comprises

for reducing the

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RESULT 8
AAU97916
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to modifying a bacterium to enhance immunogenicity of the bacterium comprising reducing the activity of an anti-apoptotic enzyme produced by the bacterium e.g. superoxide dismutase (SOD), where the bacterium has enhanced immunogenicity in a subject. Also included are the modified bacterium, an attenuated intracellular bacterium (further modified to reduce the activity of an anti-apoptotic enzyme of the bacterium), modifying a bacterium so it retains or increases immunogenicity in a subject, comprising reducing but not eliminating an activity of an enzyme produced by the bacterium, where reducing the activity of the enzyme attenuates the bacterium. The methods are useful for preventing bacterial is constitutions, e.g. tuberculosis. The attenuated intracellular bacterium is sequence is the active site of M. tuberculosis thioredoxin (TRX) which was eliminated by mutagenesis to create an attenuated bacterium of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                             Antioxidant; oxidative stress; central nervous system; cancer; ageing; neurodegenerative disorder; Parkinson's disease; Alzheimer's disease; Creutzfeldt Jakob's disease; cerebral ischaemia; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    activity of an anti-apoptotic enzyme, e.g. superoxide dismutase produced by the bacterium.
                                                                                                                                                                                                                                                                                                                                                                   motor neuron disease; scrapie; spongyform encephalopathy; acute respiratory distress syndrome; amyotrophic lateral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4 AA;
              WPI; 2002-419127/45
                                            Atlas D;
                                                                                                        26-OCT-2000; 2000GB-00026254
                                                                                                                                   26-OCT-2000; 2000GB-00026254
                                                                                                                                                                                                 GB2368339-A
                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                                                                                                             Modified-site
                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                     atherosclerosis; acquired immunodeficiency syndrome; AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tetrapeptide antioxidant peptide composition CB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU97916 standard; peptide; 4 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 CGPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                           RES
                                                                                                                                                                                                                                /note=
                                                                                                                                                                                                                                                             /note= "N terminal acetylation"
                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94; 164pp; English.
                                                                           & DEV CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 31; DB 5; 1 100.0%; Pred. No. 2e+06; tive 0; Mismatches 0;
                                                                                                                                                                                                                                តី
                                                                                                                                                                                                                                terminal amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                     sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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RESULT 9
ABB05573
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease, motor neuron diseases, scrapie, spongyform encephalopathy and loss or impaired memory, peripheral tissue disorders, particularly acute respiratory distress syndrome, amyotrophic lateral sclerosis, atherosclerotic cardiovascular disease, multiple lateral organ dysfunction, complication resulting from inflammatory processes, acquired immunodeficiency syndrome, AIDS, cancer and ageing. The antioxidant compound of the invention has the advantage that it may cross a blood barrier. The present sequence represents an antioxidant peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  may be used for preventing or reducing oxidative stress and for treating central nervous system neurodegenerative disorders, particularly Parkinson's disease, Alzheimer's disease, Creutzfeldt Jakob's disease, cerebral ischaemia, multiple sclerosis, basal ganglia degenerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to an antioxidant compound comprising a peptide including at least 3 amino acid residues of which at least 2 are cysteine residues each having a readily oxidisable sulfhydryl group for effecting antioxidation, and a first hydrophobic or non charged group attached to an amino terminal of the peptide and a second hydrophobic or non charged group attached to a carboxy terminal. the antioxidant peptide compound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antioxidant compound used for treating e.g. Alzheimer's disease and multiple sclerosis comprises peptide including cysteine residues have sulfhydryl groups and hydrophobic or non charged groups attached to
                                                                                                                                                                                                                                                                                                                                                                              Dithiol; disulfide bond; chemical reagent; sulfhydryl group; SH trans-(Pt(ethylenediamine)2Cl2)2+.
                                                                                                                                                                                                                                                                Modified-site
                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                               Dithiol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-APR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABB05573;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABB05573 standard; peptide; 4 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4 AA;
                                                                                                                                             16-MAY-2001; 2001WO-US015923
                                                                                                                                                                                                               WO200192466-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 32; 62pp; English
                                                                                                                                                                                                                                                                                               Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                             peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4; Conservative
                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                              /note= "acetylated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 31; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 5;
2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                              group;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acquired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino
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WPI; 2002-114335/15 Rabenstein DA, 26-MAY-2000; 2000US-00579883

VIND

CALIFORNIA. Shi T;

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cc amino acids derivatised for chemical peptide synthesis, where at least cone of the vials comprises an amino acid that, when fully deprotected, bears a SH group, and a vial comprising (I); (3) chemically synthesising cc a peptide comprising a disulfide linkage, involving chemically coupling a gentide comprising a form a peptide comprising at least two SH cc groups, and contacting the peptide with (I); and (4) a kit for forming an cintramolecular disulfide bond in a peptide, comprises a container containing (I). The method can be used for forming an intramolecular cc disulfide bond in a peptide. (II) is useful as a positive control when cc evaluating the efficacy of particular reagent species, and as a sample colution when optimising a purification protocol to clean up or to isolate the desired reaction product (e.g., when optimising a high cc pressure liquid chromatography (HPIC) protocol or evaluating a particular chromatography column). The present sequence represents a dithiol peptide which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a method of forming an intramolecular disulfide bond in a peptide, comprising contacting a peptide comprising at least two sulfixydryl (SH) groups with a compound (trans-
(Pt(ethylenediamine)2C12)2+) (I). Also describes are: (I) a solution (II) comprising a peptide which comprises at least two sulfur groups, and (I); a peptide synthesiser (III) for the synthesis of a peptide having an (2) a peptide synthesiser (III) for the synthesis of a vertice that in the comprises a number of vials containing intramolecular disulfide linkage, comprises a number of vials containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Forming an intramolecular disulfide bond in a peptide, comprises contacting a peptide comprising at least two sulfhydryl groups with an oxidizing agent that forms disulfide linkages in peptides.
New isolated DNA encoding a disulfide bond formation (Dsb) protein bacteria of the genus Ehrlichia, useful for preventing human monocytotropic ehrlichiosis, or canine monocytic ehrlichiosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disulphide bond formation protein; Dsb protein; gene therapy; vaccine; HME; human monocytotropic ehrlichiosis; canine monocytic ehrlichiosis; CME; antibacterial; immunostimulant; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E. coli thioredoxin active site peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-AUG-2003
                                                                                                            WPI; 2003-441356/41.
                                                                                                                                                                                                                                                             01-NOV-2001; 2001US-0335611P
                                                                                                                                                                                                                                                                                                                 01-NOV-2002; 2002WO-US035226.
                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003038061-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Escherichia coli.
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                                                                                                                                                                                                               (RERE-) RES
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Pred. No.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and polynucleotides encoding such proteins from bacteria of the genus Ebrlichia. Sequences of the invention are useful for preventing human monocytotropic ehrlichiosis (HME) or canine monocytic ehrlichiosis (CME). The kits are useful for determining whether an animal or individual has been infected with a species of the bacteria of the Ehrlichia genus. Dsb sequences are also used in gene therapy and as vaccines. The present sequence is Escherichia coll thioredoxin protein active site peptide (conserved cysteine motif) used in the exemplification of the invention
             their coding sequences, where the thioredoxin molecule is resistant to the oxidizing effects of cytokines or reactive oxygen species or resistant to S-nitrosylation of a SH-group by nitrous oxide. The mutant Trx proteins are useful for treating or diagnosing atherosclerosis, diabetes, apoptotic disease, mitochondrial dysfunction disease or cardiac dysfunction. Mitochondrial dysfunction disease include Kearns-Sayre
                                                                                                                                                                                                New mutant thioredoxin molecule which is resistant to oxidizing effects of cytokines or reactive oxygen species, useful for treating or diagnosing atherosclerosis, diabetes, Alzheimer's disease, myocardial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mitochondrial disease; cardiac dysfunction; angiogenesis dependent disease; neurodegene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antiparkinsonian; Anticonvulsant; Cardiant; Cardiovascular, Antirheumatic; Antiarthritic; Cytostatic; Antipsoriatic; Ge
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                                                                                                                                              Disclosure; SEQ ID NO 15; 145pp;
                                                                                                                                                                                                                                                                WPI; 2004-157111/15.
                                                                                                                                                                                                                                                                                                                               (UYRP)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rheumatoid arthritis; cancer; psoriasis; catalytic site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                thioredoxin; Trx; atherosclerosis; diabetes; apoptotic
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                                                                                                             present
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or encephalomyopathy
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                                                                                                                                                                                 or cancer.
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                                                                                                               invention relates
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                                                                                                elates to mutant thioredoxin (Trx) where the thioredoxin molecule is
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Pred. No. 2e+
0; Mismatches
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                                                                                                                                                English
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2e+06;
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RESULT 12
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Matches
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             The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum in the method comprises contacting the mucus or sputum of the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total volume of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian the composition further comprises incontinuate to about a statistic thorease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, multiple sclerosis or peripheral neuropathy. Cardiac dysfunction includes myocardial infarction, cardiomyopathy, arterial hypertension or heart failure. The mutant Trx proteins are also useful for treating or diagnosing angiogenesis dependent diseases, e.g. rheumatoid arthritis, cancer or psoriasis. The present sequence is the consensus sequence for the catalytic site of the Trx protein (residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Increasing the liquefaction of excessively viscous or cohesive mucus or sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucus; sputum;
lung disease; 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 15; 69pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-270016/25
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11-APR-2003; 2003US-0462082P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADM32927 standard; peptide; 4 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NAJE-) NAT
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Pred. No.
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(NADPH)
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RESULT 13
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          The present sequence is that of the antioxidant peptide, CB4. This peptide is used in a claimed pharmaceutical composition of the invention also comprising an L-DOPA amide derivative. The pharmaceutical composition is useful for treatment of a condition associated with impaired dopaminergic signalling, such as Parkinson's disease, Toursette's syndrome, attention deficit hyperactive disorder, generation of pituitar tunours and schizophrenia. The inclusion of an antioxidant in the pharmaceutical composition provides for improved stability particularly in vivo, and also provides neuroprotection against oxidative stress, which is typically associated with L-DOPA treatment, and with the biosynthesis and degradation of dopamine and other catecholamines.
                                                                                                                                                                                                                                   New L-dihydroxy phenyl alanine derivatives useful for the treatment of e.g. Parkinson's disease, Tourette's syndrome, attention deficit hyperactive disorder and schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cause of the disease. The present sequence represents an active site thioredoxin which is used in the method of the invention.
                                                                                                                                                                                                       Disclosure; Page 18; 44pp; English.
                                                                                                                                                                                                                                                                                                      WPI; 2004-625378/60
                                                                                                                                                                                                                                                                                                                                    Atlas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antioxidant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide; 4 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "C-terminal amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "N-terminal acetyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                           Tourette's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                          pituitary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of
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RESULT 14
ADV11816
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   melanoma; basal cell carcinoma; ovary tumor; breast tumor; non-small-cell lung cancer; renal cell carcinoma; bladder tumor; stomach tumor; prostatic cancer; pancreas tumor; lung tumor; uterine cervix tumor; cervical dysplasia; colon tumor; colorectal tumor; sarcoma; osteosarcoma; Kaposis sarcoma; anti-HIV; glioma; cytostatic; endocrine disease; gynecology and obstetrics; genitourinary disease; respiratory disease; metalolekeletal disease; dermatological disease; proliferative disorder; atherosclerosis; antiarteriosclerotic; cardiovascular disease; metabolic disorder; psoriasis; antipsoriatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADV11816 standard; peptide; 4 AA
              comprise a subsequence of 8 or more nucleotides or nucleotide analog wherein the subsequence is located within a sequence selected from ADV11669-ADV11724. The oligonucleotides preferably contain at least nucleotide analog, especially a locked nucleic acid (LNA) or a modi-
                                                                                                                                                                        cancer disease, or for modulating rea wive ver green boolism. proliferation, ion metabolism or glucose and energy metabolism.
                                                                                                                                                                                                         Novel compound useful for modulating expression of gene involved cancer disease, or for modulating red blood cell proliferation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immune disorder; diabetic retinopathy; antidiabetic; ophthalmological; cardiovascular disease; ocular disease; rheumatoid arthritis; antirheumatic; inflammation; asthma; antiasthmatic; anti-heumatic; inflammation; asthma; antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-MAR-2005
24-FEB-2005
                                                                                  putative human oncogene
                                                                                                  The invention relates to antisense oligonucleotides consisting of 8-50 nucleotides and/or nucleotide analogs which inhibit expression of the
                                                                                                                                                      Disclosure;
                                                                                                                                                                                                                                                                WPI; 2005-056301/06.
                                                                                                                                                                                                                                                                                               Hansen B,
                                                                                                                                                                                                                                                                                                                                                                   10-FEB-2003; 2003US-0446374P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2004241717-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cellular organisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     skin allergy; antiallergic; antiinflammatory; dermatological;
verruca vulgaris; virucide; metabolism; cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Thioredoxin conserved active site motif, SEQ ID NO:149.
                                                                                                                                                                                                                                                                                                                                    (SANT-) SANTARIS PHARMA AS.
                                                                                                                                                                                                                                                                                                                                                                                                       10-FEB-2004; 2004US-00776933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense therapy; thioredoxin inhibitor; cancer; neoplasm;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CGPC 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modulation;
                                                                                                                                                                                                                                                                                               Thrue CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                      SEQ ID NO 149; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note = Formation of this disulfide bond is reversible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 31; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell cycle; thioredoxin; cyclic.
                                                                                                                                                                                                                                                                                               Westergaard
                                                 of 8 or more nucleotides or nucleotide analogs, is located within a sequence selected from
                                                                                thioredoxin (TRX). The antisense oligonucleotides
5-methylcytosine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                               Z,
                                                                                                                                                                                                                                                                                                 Petersen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                   Wissenbach
                                                                                                                                                                                                                             involved in
                or a modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                           cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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gliomas. The TRX antisense oligonuclectides are also useful in the treatment of conditions such as atherosclerosis, psoviasis, diabetic retinopathy, rheumatoid arthritis, asthma, warts, and allergic dermatitis. They may additionally be used for inhibiting cellular proliferation and for modulating red blood cell proliferation, ion metabolism, glucose and energy metabolism, pH regulation, matrix metabolism, apoptosis, cytokinesis or the cell cycle. The TRX antisense oligonucleotides of the invention have increased specificity and affinity for TRX mRNA, and are resistant to degradation. The present sequence represents the reversibly disulfide-bonded conserved active site motif of thioredoxin which is referred to in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chemotherapeutic agent, an antiinflammatory compound or an antiviral compound. The TRX antisense oligonuclectides, and conjugates and compositions containing them, are useful in the treatment of cancers as carcinomas (e.g., malignant melanoma, basal cell carcinoma, ovarias carcinoma, breast carcinoma, non-small cell lung cancer, renal cell carcinoma, bladder carcinoma, recurrent superficial bladder cancer, stomach carcinoma, prostatic carcinoma, pancreatic carcinoma, lung carcinoma, cervical carcinoma, carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumors); sarcomas (e.g., osteosarcoma, Ewing's sarcoma, chondrosarcom
malignant fibrous histiocytoma, fibrosarcoma, and Kaposi's sarcoma);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pseudoisocytosine, 5-bromouracil, 5-propynyluracil, 6-aminopurine, aminopurine, inosine, diaminopurine and 2-chloro-6-aminopurine. The invention also relates to a conjugate comprising a TRX antisense oligonucleotide of the invention and one or more non-nucleotide or
Sequence 4 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pharmaceutical composition comprising a TRX antisense oligonucleotide conjugate of the invention, optionally further comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       origonucrectide or the invention and one or more non-nucleotide or non-polynucleotide moieties covalently attached to the oligonucleotide; and
                                                                                                  record
                                                                                                  issued
                                                                                             g
                                                                                             24-MAR-2005 : Correction to
                                                                                                  comments field
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chondrosarcoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            such
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Matches
                                                                                                           Query Match
Best Local
                 ADW97624 standard; peptide; 4 AA.
ADW97624;
                                                                          2 CGPC 5
                                                               1 CGPC 4
                                                                                                  4; Conservative
                                                                                                             Similarity
                                                                                                            100.0%;
                                                                                                            .0%; Score 31;
.0%; Pred. No.
                                                                                                   0;
                                                                                                   Mismatches
                                                                                                             2e+06;
                                                                                                                    DB
                                                                                                                    9;
                                                                                                   <u>,</u>
                                                                                                                    Length 4
                                                                                                   Indels
                                                                                                   <u>,</u>
                                                                                                   Gaps
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ADM97624
ID ADM977
XX ADM977
XX ADM977
XX ADM977
XX ADM977
XX Chron
XX Chron
XX Chron
XX Inter
KW inter
KW inter
KW inter
KW inter
XX JP200
XX JP200
XX JP200
XX JP200
XX JP200
XX IO-MA
XX IO-M chronic obstructive pulmonary disease; thioredoxin; respiratory-Gen.; antiinflammatory; cytokine antagonist; histamine antagonist; 10-MAR-2005 JP2005060408-A. Homo sapiens. chronic bronchitis. Human thioredoxin protein-related active center 19-MAY-2005 13-FEB-2004; 2004JP-00071691 30-NOV-2004; 2004JP-00346572 interleukin 2 antagonist; interleukin 4 antagonist; interleukin 6 antagonist; TNF alpha antagonist; emp (first entry)

peptide

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WPI; 2005-200153/21.

(JPCJ-) JPC KK

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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                      05-OCT-1999;
08-NOV-1999;
17-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interleukin-4-antagonists, interleukin-6-antagonists of TNP-alpha antagonists. The invention may be used for preventing and treating chronic obstructive pulmonary disease (COPD), preferably pulmonary emphysema or chronic bronchitis, caused by contaminants such as smoking gas, waste gas, dust and soot, and COPD accompanied with inflammation of cell or tissue, fibrosis and necrosis. The agent effectively prevents and treats chronic obstructive pulmonary disease, with negligible side effects. The present sequence is that of a peptide which represents the active center of thioredoxin and which is related to the
                                             New prokaryotic cell that is
status of the cytoplasm to a
disulfide bond containing pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel prophylactic or therapeutic agent of chronic obstructive pulmonary disease which contains a polypeptide of thioredoxin super family as active ingredient, preferably human thioredoxin. The invention may be used for the production of compounds
                                                                                                                                                                                                                                                                                                                                       genetic engineering; protein production; antimicrobial; antifungal; antiparasitic; antiviral; immunostimulant; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      with a respiratory-Gen. or antiinflammatory activity acting as antagonists, histamine-antagonists, interleukin-2-antagonists,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     obstructive pulmonary disease, family as active ingredient.
                          Example 6;
                                                                                                   WPI; 2005-239572/25.
                                                                                                                         Beckwith J,
                                                                                                                                                   (HARD
                                                                                                                                                                                                                                        05-OCT-2000;
                                                                                                                                                                                                                                                                 29-MAR-2005
                                                                                                                                                                                                                                                                                        US6872563-B1
                                                                                                                                                                                                                                                                                                                                                                             Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                      02-JUN-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADY81685 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prophylactic/therapeutic agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Agent e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N
                                                                                                                                                  HARVARD COI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .g. tablet or capsules,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     - CGPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGPC
                          SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ
                                                                                                                                                                                                                                                                                                                                                                            TrxA wild type redox active disulfide motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                         Aslund F,
                                                                                                                                                                                                                                        2000US-00679705
                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ທ
                                                                                                                                                                                     99US-0157770P.
99US-0163939P.
99US-0166044P.
                          ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              占
                                                                                                                                                                COLLEGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ŏ
                                                                                                                                                   SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                          2;
                        44pp; English
                                                                                                                          Bessette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10pp;
                                                 proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>.</u>
                                                             genetically more oxidati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for preventing and treating chronic contains polypeptide of thioredoxin super
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                          PH,
                                                             tically modified oxidative state,
                                                                                                                        Georgiou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 4;
                                                             to shift, useful i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                          Ritz
                                                                                                                         á
                                                                                                                                                                                                                                                                                                                                                                             SEQ ID
                                                              for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   as cytokine-
                                                             e redox
producing
                                                                                                                                                                                                                                                                                                                                                                               NO: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                          JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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invention relates

to a

prokaryotic cell that is genetically modified

This sequence represents a conserved active site found in thioredoxins. The thioredoxins are a class of low molecular weight redox proteins. Thioredoxin exerts specific redox control over a number of transcription

DNA binding and therefore regulates

to modulate

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RESULT 17
AAY78911
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S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            favors disulfide bond formation, relative to a prokaryotic cell that is not genetically modified. The cell further comprises a mutated AhpC gene comprising an insertion of three nucleotides in the TCT triplet rich region located at codons 36-9 of an AhpC gene. The insertion increases the cell's ability to proliferate relative to a cell that is not further genetically modified. Also described is a method for producing a protein having at least one disulfide bond. The prokaryotic cell and method are useful for producing disulfide bond containing proteins. The proteins, polypeptides, and compositions are useful for diagnostic, therapeutic, or prophylactic purposes. They are useful for inhibiting the growth of bacteria, viruses, fungi, or other parasites and for stimulating an immune response. They may be used as food supplements, as well as
                                                                                                                                                                                                                                                                                                                                                                                                                                                             wound healing; Crohn's disease; ulcerative colitis; angiogenesis; adult respiratory distress syndrome; ARDS; infection; inflammatory conditions, psoriasis; inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             apoptosis; reperfusion injury; hepatitis; amyotrophic lateral scler Alzheimer's disease; autoimmune disease; Sjogren's syndrome; lupus; rheumatoid arthritis; HIV; Hermansky-Pudlack syndrome; retinopathy;
                                                                                                                         New composition for preventing inhibition of apoptosis and for treating e.g. cancer or reperfusion injuries comprises inhibitor of cellular redox
                                                                                                                                                                                                      Kirkpatrick DL,
                                                                                                                                                                                                                                                                                    31-ЛЛТ-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thioredoxin reductase; redox inhibitor; cellular proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 4 AA;
                                                                                                                                                                          WPI; 2000-195158/17.
                                                                                                                                                                                                                                                                                                                    02-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                   WO200006088-A2
                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                               hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY78911 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   components of wash
                                                                                                                                                                                                                                                                                                                                                    10-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               o shift the redox status of the cytoplasm to a more oxidative state that
                                                                                                                                                                                                                                      (KIRK/) KIRKPATRICK D
(POWI/) POWIS G.
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                                                                            Page 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               conserved active site amino acid sequence.
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redox active disulfide motif.
                                                                                                                                                                                                      Powis
                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease;
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Pred. No. 2
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RESULT 18
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12-JUL-1991;
19-AUG-1991;
01-NOV-1991;
                                                                                                                                                                                                New T cell-derived colony stimulating factor for treating compromised patients, including acquired immunodeficiency patients and certain types of cancer patients.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T cell-derived colony stimulating factor; TC-CSF; immunostimulant; acquired immune deficiency syndrome; AIDS; cancer; gene therapy; haematopoietic growth factor; human; thioredoxin; ADF; active site
The invention relates to T-cell derived colony stimulating factors (TC-CSF) and nucleic acid molecules encoding them. TC-CSF which is a novel haematopoietic growth factor is useful for treating immune-compromised
                                                                                                                                                                                                                                                                                                                                             WPI; 2001-432041/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-JUL-2001
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                                                                                                                                            Example 33;
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                                                                                                                                      Col 36; 38pp; English
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91US-00729135.
91US-00747784.
91US-00788115.
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Matches 4
                                                                                                                  sequence
relating
                                                                                                                          The invention relates to the use of thioredoxin as a cell growth stimulator, for modulating thioredoxin cell growth stimulator-associated inhibition of apoptosis, comprising controlling the expression and/or activity of the thioredoxin. The invention is useful for treating tumours associated with aberrant thioredoxin expression and activity. This sequence represents a wild type thioredoxin active site peptide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            patients, including acquired immune deficiency syndrome (AIDS) patients and certain types of cancer patients. TC-CSF DNA is used in gene therap. The present sequence is human thioredoxin active site which is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; thioredoxin; cell growth stimulator; modulating; apoptosis; tumour; wild type.
                                                                                                                                                                                                                                                                                                                                           03-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wild type thioredoxin active site peptide sequence
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                                                                                          Sequence 5
                                                                                                                                                                                                              Disclosure; Page 1; 59pp; English.
                                                                                                                                                                                                                                     treatment of tumors.
                                                                                                                                                                                                                                                Modulating thioredoxin-associated inhibition of apoptosis for the
                                                                                                                                                                                                                                                                      WPI; 2002-470971/50
                                                                                                                                                                                                                                                                                               Powig
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAO21523;
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                                                                                                                                                                                                                                                                                                                     (POWI/) POWIS
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tive 0; Mismatches
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Pred. No. 2e+
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                                               Gaps
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ABG71301 ID ABG XX

ABG71301 standard; peptide; 5 AA

RESULT 20

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The invention relates to isolated human thioredoxin polypeptide TRXP-2, a naturally occurring polypeptide comprising an amino acid sequence 90% comparison and the sequence of TRXP-2 and its encoding polymucleotide (TRXP-2 NA). Also included are a recombinant polymucleotide comprising a promoter sequence operably linked to TRXP-2 NA, a cell transformed with the recombinant sequence, a TRXP-2 ctransgenic organism, an anti-TRXP-2 antibody and a composition comprising compound identified using TRXP-2, an agonist or antagonist compound identified using TRXP-2, or the close antibody. TRXP-2 is useful for screening a compound for condition associated with decreased or overexpression of functional TRXP cin a patient. TRXP-2 is useful for screening a compound that condition associated with decreased or overexpression of functional TRXP cin a patient. TRXP-2 is also useful for screening for a compound that condition at the condition of the polypeptide of the sequence of TRXP-2 NA is consisted if or screening a compound that condition of a target polymucleotide comprising the sequence of TRXP-2 NA. The contibody is useful for detecting the presence of TRXP-2 NA. The contibody is useful for detecting the presence of TRXP-2 NA. The contibody is useful for detecting the presence of TRXP-2 NA and modulators of conditions or disease associated with expression of TRXP in a subject or in a biological sample. TRXP-2 and TRXP-2 NA and modulators of conditional prevention of cell conditional vera, patient, mixed connective tissue confidence, polycythaemia vera, patiency distress syndrome, Addison's confidence of languages. The antibody of the polycome, and prevention of cell inflammation, osteoprosis, reumatoid arthritis. Reiter's confidence and prevention or confidence and preventions.
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scleroderma, systemic lupus erythematosus, ulcerative colitis, Reiter's syndrome, haemodialysis, Crohn's disease, atopic dermatitis, autoimmune thyroiditis, diabetes mellitus, Graves' disease and glomerulonephritis)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel human thioredoxin protein and polynucleotide for diagnosing, preventing and treating cell proliferative, inflammatory and viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-2001; 2001US-00954846
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer; actinic keratosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Guegler KJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        atherosclerosis; bursitis;
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Sequence 6

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RESULT 21
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ID AAP71
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AC AAP71
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Best Local S
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31-OCT-2002
06-JUN-1991
                            It is a redox active peptide which can be used in cpds. to treat stress and ischaemia, for inhibiting lipid peroxidin. due to drugs or toxic effects, as anti-inflammatory agents, to prevent DNA damage, and to maintain the biological activity of antibiotics or proteins. Thioredoxin can be used as a general antioxidant in the food and cosmetic industry. It may be used in prodn. processes for antibiotics, proteins or polypeptides. Therapeutic dose is 1 microgram/kg - 100mg/kg. (Updated on 25-WAR-2003 to correct
                                                                                                                                                                    New and known thioredoxin derivs. and analogues - useful for metal catalysed oxidative damage in biological reactions and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infection (e.g. viral, bacterial, fungal, parasitic, protozoal an helminthic), viral disorders (e.g. acquired immunodeficiency synd (AIDS) and trauma. The present sequence represents a thioredoxin site disulphide/dithiol conserved sequence as found in TRXP-2
                                                                                                                                                                                                                            Pigiet
                                                                                                                                                                                                                                                                        14-MAR-1986;
20-OCT-1986;
                                                                                                                                                                                                                                                                                              10-FEB-1986;
                                                                                                                                                                                                                                                                                                                      10-FEB-1987;
                                                                                                                                                                                                                                                                                                                                            16-SEP-1987
                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Redox active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence of thioredoxin analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAP71087;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAP71087 standard;
                                                                                                                                                                                                      WPI; 1987-258442/37.
                                                                                                                                                                                                                                                                                                                                                                 EP237189-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            food industry;
                                                                                                                                                                                                                                                  (REPK ) REPLIGEN CORP.
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                                                                                                                                  Page 7; 7pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGPC 5
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                                                                                                                                                                                                                            Millis CD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide; antioxidant; stress; ischaemia; lipid peroxidation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
                                                                                                                                                                                                                                                                       86US-00828112.
86US-00839857.
86US-00921287.
                                                                                                                                                                                                                                                                                                                     87EP-00301150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cosmetics; antibiotic
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terminal blocking
                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                         /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                  English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .0%; Score 31;
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                                                                                                                                                                     disease
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RESULT 22
ABP54936
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                                                                                                                           The present sequence is that of a hexapeptide comprising the active site sequence of thioredoxin. The hexapeptide was used in an example from the invention that examined the binding of 4-(N-(S-glutathionylacetyl)amino)phenylarsenoxide (GSAO) to peptide and protein dithiols. The results indicated that GSAO selectively binds proteins containing closely spaced thiols. The invention provides a method for the treatment and/or prophylaxis of arthritis using a compound of formula A-(I-Y)p, where A is a cell-membrane impermeable pendant group, L is a clinker and/or spacer, Y is an areenoxide or arsenoxide equivalent, and p is an integer from 1 to 10. The compound, e.g. GSAO, is useful for the treatment of arthritic conditions such as calcific periarrhritis, contained, septic or Jaccoud's arthritis, gout, hand or knee osteoarthritis, of guerinitis, and polyarthritis (all claimed). The compound blocks angiogenesis in the synovial tissue and leucocyte ingress that triggers inflammation, with no signs or symptoms of toxicity
                                                  Query Match
Best Local S
Matches 4
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Best Local S
Matches 4
                                                                                                       Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                           arthritis in
                                                                                                                                                                                                                                                                                                                                                                                                       Use of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-MAR-2001; 2001AU-00003798.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200274305-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABP54936;
                                                                                                                                                                                                                                                                                                                                                               Example 2(b); Page 42; 91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-MAR-2002; 2002WO-AU000310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Active site peptide of thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABP54936 standard; peptide; 6 AA.
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                                                  Similarity 4; Conserv
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                         CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                       impermeable cell membrane compound
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                                                                                                         B,
                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  organoarsenical; arsenoxide; antiarthritic; antirheumatic; arthritis; therapy.
                                                                                                                                                                                                                                                                                                                                                                                           vertebrates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
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                                                               100.0%; Score 31;
100.0%; Pred. No.
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Pred. No.
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                                                   Mismatches
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2e+06;
                                                               2e+06;
                                                                             DB 5; Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                      for the treatment
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                                                   Indels
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RESULT

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RESULT 24
ADH05024
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DT 11-MP
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Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a method for producing an antiangiogenesis plasmin fragment. The method comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin reductase, where a reduced plasmin protein is produced and the antiangiogenesis plasmin fragment having anti-angiogenesis activity, is released from the reduced plasmin protein. The plasminogen activator is preferably a urokinase-type plasminogen activator. The angiogenesis plasmin fragment is A61. The plasmin reductase is selected from annexin II heterotetramer, annexin II p36 subunit, p11, thioredoxin, and protein disulphide isomerase. The annexin II heterotetramer is associated with a cell membrane. The method of the invention is useful for modulating (e.g. promoting or inhibiting) angiogenesis. The present sequence represents a mammalian cleavage peptide found in thioredoxin and protein disulphide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  plasmin reductase; reduced plasmin protein; plasminogen activato urokinase-type plasminogen activator; A6; annexin II heterotetramer; annexin II p36 subunit; annexin II p11 subunit; thioredoxin; protein disulphide isomerase; modulating angiogenesis; mammalian; cleavage peptide.
11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Producing an anti-angiogenesis plasmin fragment, useful for modulating, e.g. promoting or inhibiting angiogenesis, comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin
                                                                 ADH05024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-596985/56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Waisman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-NOV-2002; 2002US-00304287.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAY-2003
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                                                                                                                         ADH05024 standard; peptide; 6 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kwon M;
                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peptide; 6 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 31; DB 6;
100.0%; Pred. No. 2e+06;
live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 6;
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RESULT 25
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Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a molecule with two pendant phenylarsine moieties. The invention is useful for imparting detectable properties to target material, for detecting target material, for monitoring binding process, for isolating and immobilizing at least one target material, for screening a library of peptides and identifying a cysteine containing peptides and for labeling protein in living cells. The present sequence represents a target peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New bis-phenylarsine derivatives useful for labeling protein in living cells and for screening a library of peptides and identifying a cysteine
                                                                                           Modified-site
                                                                                                                                                     Synthetic
                                                                                                                                                                                         mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive sputum;
                                                                                                                                                                                                                                         17-JUN-2004
                                                                                                                                                                                                                                                                ADM32929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-JUN-2002; 2002US-0388699P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-JUN-2003; 2003WO-US018792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pendant phenylarsine moieties; labeling; target peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Target peptide
                                                                                                                   Active-site
                                                                                                                                                                                                               Amino acid sequence
                                                                                                                                                                                                                                                                                       ADM32929 standard; peptide; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ebright RH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003107010-A1
                                   WO2004024868-A2
                                                                      Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RUTF ) UNIV RUTGERS STATE NEW JERSEY
                                                                                                                                                                             disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2004-156348/15.
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                                                                                                                                                                                                                                                                                                                                                                            2 CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                  4.
                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 5; 82pp; English
                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ebright YW,
                                                                                                                                                                                                                                                                                                                                                    ហ
                                                                                                                                                                            cystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 #5 of the invention
                                                                                                                               Location/Qualifiers
                                                                                /note= "this residue is in a reduced state"
                                                         note= "this residue is"
                                                                                                        'note= "active site of thioredoxin"
                                                                                                                                                                                                                 of the thioredoxin active site
                                                                                                                                                                                                                                       entry)
                                                                                                                                                                             fibrosis.
                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 31; DB 8; 100.0%; Pred. No. 2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                        in a reduced state
                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                         Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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25-MAR-2004

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a tott volume of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin. The composition further comprises nicotinamide-adenine thioredoxin. The composition further comprises nicotinamide-adenine dinucleotide phosphate (reduced from) (NADPH) for reducing the thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesiveness or sputum. These may be used for treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputum is a symptom or consistency or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputum is a symptom or cohesiveness of mucus or sputu
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6 AA;
                       Misc-difference
                                                                                                                                           Synthetic.
                                                                                                                                                                                                                      mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive sputum;
                                                                                                                                                                                                                                                                        Amino acid
                                                                                                                                                                                                                                                                                                                      17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                       ADM32928;
                                                                                                                                                                                                                                                                                                                                                                                                                      ADM32928 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cause of the disease. The present sequence represents an thioredoxin which is used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum. The method comprises contacting the mucus or sputum of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 13; Page 51; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Increasing the liquefaction of excessively viscous or cohesive mucus sputum comprises contacting the mucus or sputum with a composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2004-270016/25
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                                                                     Active-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprising a protein or peptide containing a thioredoxin active-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-SEP-2002;
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                                                                                                                                                                                            lung disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CGPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CGPC 5
                                                                                                                                                                                                                                                                        sequence of the thioredoxin active site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JEWISH MEDICAL & RES CENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2003WO-US028526
                                                                                                                                                                                                                                                                                                                    (first entry)
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2003US-0462082P.
                                                                                                                                                                                            cystic
                                                                                          Location/Qualifiers
/note= "any amino acid"
                                              note= "active site of thioredoxin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      describes a method for increasing the liquefaction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             represents an active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8; Length 6;
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RESULT 27
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                              statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian thioredoxin. The composition further comprises nicotinamide-adenine dinucleotide phosphate (reduced form) (NADPH) for reducing the thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for the protein of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total volume of a sample of mucus or sputum from the patient shows a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum. The method comprises contacting the mucus or sputum of
                                                                                                                                                                                                                                                                                                                              treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. The present sequence represents an active site of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Increasing the liquefaction of excessively viscous or cohesive mucus or sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
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ADP87509;
                                    ADP87509 standard; peptide; 6 AA.
                                                                                                                                                                                                                                                                           Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 12; Page 51; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reduced state.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-SEP-2002;
11-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-SEP-2003; 2003WO-US028526
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                                                                                                                                                                                                                                                                                                               thioredoxin which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NAJE-) NAT JEWISH MEDICAL & RES CENT
                                                                                                                             N
                                                                                                                                                                2 CGPC 5
                                                                                                                                                                                                  4.
                                                                                                                                                                                                                       Similarity
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2003US-0462082P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "this residue is in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "any
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "this residue is in a reduced state"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "any amino acid'
                                                                                                                                                                                                                                                                                                               is used
                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                             The present sequence represents an active site of a used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino
                                                                                                                                                                                                    0
                                                                                                                                                                                                                     Score 31; DB 8;
Pred. No. 2e+06;
                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        reduced state"
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                                                                                                                                                                                                                                     Length 6;
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RESULT 28
ADP67423
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                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                  cyclohexa-2,5-dienone derivatives (I) and their salts, esters, anides, and the setters, anides, and can be used as thioredoxin reductase inhibitors. (I) can be used for treating a proliferative condition (especially colon or renal cancer) or a condition mediated by thioredoxin/thioredoxin reductase in the human or animal body. (I) are also useful for inhibiting thioredoxin/thioredoxin reductase and call cycle progression, to regulate/inhibit cell proliferation and/or to promote apoptosis in a cell in vitro or in vivo. The present sequence represents a thioredoxin family conserved catalytic site peptide, which is given in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2; 5-dienone;
cytostatic; thioredoxin reductase inhibitor; proliferative condition;
colon cancer; renal cancer; thioredoxin; thioredoxin reductase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New 4-(1(sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2,5-dienone derivatives are thioredoxin reductase inhibitors useful for treating e.g. colon or renal cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell cycle progression; cell proliferation; apoptosis; catalytic site peptide.
                                                                                                                                                                                                                                                                                                                                                                   Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-DEC-2002; 2002WO-GB005842
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                                                    p11; p11 activity modulator; plasminogen activation; cytostatic;
antisense therapy; cancer; tumour; tumour growth inhibition; thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 82; 141pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-517586/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stevens
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                                       protein disulfide
                                                                                               Thioredoxin
                                                                                                                          23-SEP-2004
                                                                                                                                                      ADP67423;
                                                                                                                                                                                  ADP67423 standard; peptide; 6 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention described 4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)-5-dienone derivatives (I) and their salts, esters, amides,
                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                             protein disulfide isomerase common
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              conserved catalytic site peptide.
                                                                                                                                                                                                                                                                                                                         100.0%; Score 31; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                               Mismatches
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Synthetic

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RESULT 29
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XX ADQ97
XC ADQ97
XX 07-00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a composition (I) which modulates the CC activity of a p11 protein and effects a change in the level of CC plasminogen activation by a cell. Also described: (1) making (M1) a CC clonal cell line, which involves isolating a cell, then characterising CC the activity of a protein produced by the cell or clonal progeny of the CC cell, where the protein is involved in plasminogen activation; and (2) a CC colonal cell line (II) useful in the identification of composition that CC modulate p11 activity, where the clonal cell line is obtained by (M1). (C) has cytostatic activity, and can be used in antisense therapy. (I) is useful for modulating the activity of p11 which involves administering (I) to a cell. The cell can be a human cancer cell, chosen from a HT1080 (I) to a cell. The cell can be a human cancer cell, chosen from a HT1080 (C) tibrosarcoma cell, a LNCap prostate cancer cell and a CCI-22 colorectal adenocarcinoma cell, a LNCap prostate cancer reducing the development of cancer cell in the patient. (I) is useful for reducing the growth of CC cancer cell in the patient. (I) is useful for inhibiting the growth of CC cancer cell in the patient. (II) is useful for composition that modulates p11 activity which involves administering the composition that modulates p11 activity which involves cadministering the composition that modulates p11 activity which involves call of a clonal cell line that had not received the composition, and cclaim is composition that produces a change in p11 activity. The composition that produces a change in p11 activity. The composition sequence represents a thioredoxin and protein confidentification sequence represents a thioredoxin and protein confidentification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Composition useful for inhibiting growth of tumor in patient, modulates activity of pl1 protein and effects change in level of plasminogen activation by a cell.
                                         pll protein; plasminogen activation; tumour growth; metastasis;
                                                                                  Thioredoxin
                                                                                                                            07-OCT-2004
                                                                                                                                                                                                          ADQ97008 standard; peptide; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-DEC-2002; 2002US-0433140P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-DEC-2003; 2003WO-US040029
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-487995/46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MEDI-) MEDIOMICS
(WAIS/) WAISMAN D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present
                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIOMICS LLC
                                                                                                                                                                                                                                                                                                                                                        CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A,
                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 31; DB 8; Length 6; ilarity 100.0%; Pred. No. 2e+06; Conservative 0; Mismatches 0; Indels
                                                                                  and protein disulphide isomerase common
                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention.
                                                                                sequence.
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a composition which modulates the activity of a pli protein and effects a change in the level of plasminogen activation by a cell. The composition is useful for reducing pli protein activity in cancer cells. They are also useful for inhibiting tumour growth or metastasis or treating cancer. The present sequence represents the amino acid sequence of the thioredoxin and protein disulphide isomerase common sequence involved with cleavage, formation and reshuffling of disulphide
                                                                                                                                                                                                                                               Sequence of oligopeptide 3 specific to the beta-subunit of luteinising hormone (\operatorname{LH}) .
                                                                                                                                                                                                                                                                                                                                    AAP50203 standard; peptide; 8 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New composition which modulates the activity of a pll protein and effects a change in the level of plasminogen activation by a cell useful for treating cancer or reducing pll protein activity in cancer cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-NOV-2001; 2001US-0333866P.
26-NOV-2002; 2002US-00304287
13-DEC-2002; 2002US-0433140P
                                   Maes RF;
                                                                                                                                   22-MAY-1985.
                                                                                                                                                                                   Ното варіелв
                                                                                                                                                                                                           Contraception; assay; pregnancy test; luteinising hormone; human chorionic gonadotropin.
                                                                                                                                                                                                                                                                                                             AAP50203;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; SEQ ID NO 4; 66pp; English.
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                                                                                   26-AUG-1983;
                                                                                                            23-AUG-1984;
                                                                                                                                                                                                                                                                                     17-OCT-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Waisman DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-DEC-2003; 2003US-00735577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (WAIS/) WAISMAN D M.
                                                          (ANDA-)
                                                                                                                                                                                                                                                                                                                                                                                                   2 CGPC 5
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                                                           ANDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                  4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                          BIOLOGICALS
                                                                                                                                                                                                                                                                                  (first entry)
                                                                                   83EP-00401714.
                                                                                                            84EP-00401710
                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 31; DB 8;
Pred. No. 2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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WPI; 1985-124152/21

WPI; 1995-060814/08

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RESULT 31
ADC811369
ID ADC8137
AC ADC81
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HCG, located near cysteine groups, are used in a claimed process for the preparation of anti-HCG and anti-LH vaccines. The vaccines are characterised in that these oligopeptides are complexed upon themselves and between themselves by disulphur bridges, upon their own beta-subunit of HCG and LH or upon oligopeptides containing cysteine whose AA chain is common to LH and HCG, namely the peptides in AAP50208-P50212. Through sulphur bridges the specific oligopeptides are administered by injection in vaccines to cause prodn. of antibodies so that a contraceptive effect in vaccines to cause prodn. of antibodies so that a contraceptive effect.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Con A-induced interleukin-2 production; mouse spleen cells; granulocyte-monocyte colony forming cell; GPCC; mouse bone marrow; immunomodulatory agent; congenital immune defect; primary T cell deficiency; thymic dysplasia; Down's syndrome; viral infection; herpes; adenovirus infection; HV infection; subscute sclerosing pan-encephalitis; immune suppression; leukopaenia; cancer; chemotherapy; radiotherapy; autoimmune inflammatory disorder; rheumaroid arthrifis; systemic lupus exythematosus; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligopeptides AAP50198-P50205, specific to the beta-subunit of LH and HCG, located near cysteine groups, are used in a claimed process for t preparation of anti-HCG and anti-LH vaccines. The vaccines are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prepn. of anti-HCG and anti-LH vaccines - by use of oligo-peptide(s) specific to beta-sub-unit of LH and HCG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 8 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 17; 22pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bovin THF-gamma2 analogue C3-C6-cyclicTHF gamma2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JAN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADC81369 standard; peptide; 8 AA
                                   Rycus
                                                                                             (YEDA ) YEDA RES & DEV CO LTD.
                                                                                                                                                                                                                          28-JUN-1994;
                                                                                                                                                                                                                                                                                    12-JAN-1995
                                                                                                                                                                                                                                                                                                                                                 WO9501182-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mutant; mutein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bone marrow transplantation; atopic condition; asthma; atopic dermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cow; thymic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              achieved; and the body fluids
                                   P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 CGPC 5
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4; Conservative
                                      Burstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               humoral factor gamma2; THF-gamma2; concanavalin
                                                                                                                                                                                                                          94WO-US007304.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers 3. .6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                          /note= "Wild-type Lys substituted by Cys'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Wild-type Asp substituted by Cys"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 note= "This bond cycylises the molecule"
                            Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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                                      Trainin N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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Sequence 8 AA;

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CC free carboxyl group and a free amino group or through a cysteine residues or (v) linkage of two or more sequences or modified CC sequences corresponding to (i)-(iv), either directly or through a spetide cornon-peptide chain. The peptides are used as immune defects involving CC primary T cell deficiencies such as thymic dysplasia and Down's syndrome, CC primary and secondary viral infections (e.g. herpes virus, adenovirus and CC HIV), as well as subacute infections such as subacute sclerosing pan-CC encephalitis, immune suppression and leukopaenia following cancer treatment by chemotherapy and/or radiotherapy, autoimmune inflammatory disorders, e.g. rheumatoid arthritis, systemic lupus exythematosus and CC psoriasis, in bone marrow transplantarion to prevent viral infections and CC in atopic conditions such as asthma and atopic dermatitis. The present CC sequence is a bovine THF-gamma2 analogue peptide of the invention
A peptide is claimed which is a thymic humoral factor (THF)-gamma2 analogue of at least 4 amino acid (AA) residues or a functional derivative or salt, capable of enhancing concanavalin A (Con A)-induction the releukin-2 (IL-2) production in mouse spleen cells and/or the numb of granulocyte-monocyte colony forming cells (GM-CFC) of mouse bone marrow. The peptide comprises one or more sequences corresponding to sequence of the wild-type THF-gamma2 but differing by (i) deletion of or more AA residues, (ii) addition of one or more AA residues by a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural AA residue, (iv) cyclisation through a protein natural or non-natural capacity cyclisation through a protein natural capacity ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New thymic humoral factor gamma 2 analogues - used as immunomodulatory agents for treating e.g. immune defects, viral infections or auto:immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 17; 30pp; English
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AAW05002
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Best Local :
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                                                                                                 Immunomodulation; immunomodulator; immune system; anergy; deficiency;
aberrant response; chronic bronchitis; atopic disease; AIDS;
acquired immune deficiency syndrome; herpes virus infection; cyclic.
                                                                                                                                                            Thymic humoral
                                                                                                                                                                                         03-DEC-1996
                                                                                                                                                                                                                                                AAW05002 standard; peptide; 8
                                                                      Synthetic
                                                                                                                                                                                                                     AAW05002;
WO9611943-A1
                           Disulfide-bond
                                                                                                                                                                                                                                                                                                                        w
                                                                                                                                                                                                                                                                                                                                                   2 CGPC 5
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                                                                                                : response;
                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                        (first entry)
                                                                                                                                                           factor gamma-2 cyclic analogue.
                           Location/Qualifiers
3. .6
                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                               Score 31; DB
Pred. No. 2e+
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                DB 2;
2e+06;
                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                             Length 8;
                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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06-OCT-1995;

95WO-SE001151

25-APR-1996.

14-OCT-1994;

94SE-00003526

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RESULT 33
AAW41664
ID AAW41
XX AAW41
AC AAW41
AC
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Best Local S
Matches 4
A novel immunosuppressing or immunostimulatory peptide conforms to the motifs represented by the formulae of the invention, with the proviso that the peptides described in AAW36999, AAW37000 and AAW41647 to AAW41682 are excluded. An immunosuppressant can be used to treat transplant rejection or autoimmune disease, e.g. rheumatoid arthritis, systemic lupus erythematosis, Sjogren's syndrome, scleroderma, mixed connective tissue disease, dermatomyositis, polymyositis, Reiter's syndrome, Behcet's disease, type I diabetes, Hashimoto's thyroiditis,
                                                                                                                                                                                                                                                                                                                                            Nucleic acids encoding cysteine- or methionine-containing peptide(s) which have immuno:stimulatory or immunosuppressive activity - can be to treat, e.g. cancers, infection, auto:immune disease or transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present peptide is a known immunomodulatory agent. It is described as a cyclic analogue of thymic humoral factor gamma-2 and was disclosed in WO9501182. The peptide is one of 6 known peptides which concord with a highly generic sequence covering immunomodulatory peptides of 4-15 amino acids being claimed in the present patent application; all 6 peptides are specifically excluded from the claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide(s) with immunomodulatory activity - useful for treating e chronic bronchitis, malignancies, herpes, AIDS or atopic disease, adjuvants in vaccines, or for inhibiting graft rejection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunomodulator;
transplant reject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 8 AA;
                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bergstrand H,
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23-SEP-1996;
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96SE-00003469
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                                                                                                                                                                                                                                                           19; 183pp; English.
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ion; autoimmune disease; cancer; infi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide; 8 AA.
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Graves' disease, multiple sclerosis, myasthenia gravis, encephalomyelitis, phemphigus vulgaris, vegetans or foliaceus, Usher syndrome or Brazilian phemphigus. An immunostimulator cau to treat conditions such as cancer or infection
                                                                                                                         The present invention relates to a molecule with two pendant phenylarsine molecies. The invention is useful for imparting detectable properties to target material, for detecting target material, for monitoring binding process, for isolating and immobilizing at least one target material, for screening a library of peptides and identifying a cysteine containing peptides and for labeling protein in living cells. The present sequence
                                                                                        Sequence 8 AA;
                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                           New bis-phenylarsine derivatives useful for labeling protein in living cells and for screening a library of peptides and identifying a cysteine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Target peptide
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                                                                                                                                                                                                                              containing peptides
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pendant phenylarsine
                                                                                                                 represents a target peptide of the invention.
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                                                                                                                                                                                                         25; SEQ ID NO 10; 82pp; English.
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100.0%; 1
tive 0;
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Pred. No. 2e+
0; Mismatches
                                             Score 31; DB
Pred. No. 2e+
0; Mismatches
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2e+06;
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                                                                Length 8;
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an be use
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RESULT 35 ADV60194 ID ADV60 XX

ADV60194

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis; cancer; obesity; diabetes; asthma; inflammation; de food; feedstuff; cosmetics; agriculture; animal breeding; VEGF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-MAR-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the family of protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUN-2004; 2004WO-EP051158.
                ADZ28911;
                                                                                                                                                                                                                                                                                  Sequence 8
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17-JUN-2003; 2003US-0479061P.
                                                  ADZ28911 standard; protein; 8
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100.0%; Pred. No. 2e+06;
:ive 0; Mismatches
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RESULT 37
ADV60195
ID ADV60

ADV60195 standard; peptide; 9 AA

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Matches Query Match

4; Conservative Similarity

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Mismatches

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Gaps

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100.0%; Score 31; DB 9; 100.0%; Pred. No. 2e+06;

DB 9;

Length 8;

Local

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The invention relates to a cellular extract from an organism comprising a CC SlyD gene, substantially free of a SlyD polypeptide that binds to a bi-CC arsenical reagent. Also described are: (i) a kit for the in vitro CC production of proteins, the kit comprising a cellular extract above; (ii) a polypeptide comprising an amino acid sequence that binds specifically to EDT 2 [4',5'.bis(1,3,2-dithioarsolan-2-yl) fluorescein-(1,2-ethanedithiol)2. Preferably, the cellular extract is cc entered that binds specifically to EDT 2 [5'.bis(1,3,2-dithioarsolan-2-yl) fluorescein-(1,2-ethanedithiol)2. Preferably, the cellular extract is cc m Escherichia coli extract or from an E. coli comprising an amino acid complete coli extract is specifically to EDT 2 [5'.bis(1,3,2-dithioarsolan-2-yl) fluorescein-(1,2-ethanedithiol)2. Preferably, the cellular extract is cellular extract is sequence that binds a bi-arsenical molecule. The SlyD mutant gene encodes a SlyD that is mutated in an amino caid sequence that binds a bi-arsenical molecule. The SlyD mutant gene elso encodes a truncated slyD protein. The cellular extract has reduced activity of at least one enzyme that catalyzes hydrolysis of high-energy phosphate bonds or hydrolysis or formation of phosphodiester bonds. The cellular extract has that catalyzes hydrolysis or hydrolysis or formation of phosphodiester bonds. The cellular extract further comprises a nuclease sproviding chemical energy for synthesis. It also comprises a nucleic acid encoding a fusion protein encoding an cc exogenous protein fused to a tag, where the tag binds a bi-arsenical cc exogenous protein for purifying and detecting recombinant polypeptides chaving polyhistidine or polycysteine tags. The present sequence represents calmodulin ortho c-terminus used as a control in measurement cc of represents entered in vitro protein synthesis.
  New cellular extract from an organism comprises a SlyD gene, substantially free of a SlyD polypeptide that binds to a bi-arsenical reagent, useful for synthesizing, purifying, or detecting biomolecules,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell culture; DNA calmodulin ortho;
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Sequence 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kudlicki WA,
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RESULT 38
AAR50846
ID AAR50
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AC AAR50
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                                                                                                                                                                                                                                                                                           genetic diseases such as cancer, obesity, diabetes, asthma and configuration, neuropsychiatric disorders such as depression, for quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway canalysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present of sequence represents a peptide from a VEGF peptide combo.
                                                                                                                                                                                             Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex
   AAR50846;
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                                 AAR50846 standard; peptide; 10
                                                                                                                                                                                                                                                                 Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JUN-2003; 2003EP-00101775.
17-JUN-2003; 2003US-0479061P.
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4; Conserv
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The peptide modifies the binding of P-, E- and L-selectin and of integrins, especially intercellular adhesion molecule-1 (ICAM-1). It inhibits cellular adhesion. It can be used to treat or prevent inflammation, coagulation, ischaemia, reperfusion injury, bacterial sepsis, disseminated intravascular coagulation, adult respiratory distress syndrome, tumour metastisis, rheumatoid arthritis and atherosclerosis. When labelled it can be used to detect defective selectin (or integrin) binding ligands or high (localised) concentrations of selectin (or integrin) binding cells, especially leucocytes or tumour cells. It can also be used for in vivo imaging. This is a specific example of a degenerate sequence given in the specification. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Selectin; integrin; intercellular adhesion molecule; ICAM-1; inh inflammation; coagulation; ischaemia; reperfusion injury; bacterial sepsis; tumour metastisis; bacterial sepsis; tumour metastisis; disseminated intravascular coagulation; rheumatoid arthritis; adult respiratory distress syndrome; atherosclerosis; treatment; prevention; detection; tumour; leucocyte; imaging.
Sequence 10
                                                                                                                                                                                                                                             New cyclic peptide(s) inhibiting selections and integrins - preven cellular adhesion in treatment of inflammation, perfusion, injury, metastasis, etc. also for diagnosis.
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27-SEP-1994
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(first entry)
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Query Match

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Score

31; DB 2;

Length 10;

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RESULT 39
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Matches
                                                                                                    New cyclic peptide(s) inhibiting selections and integrins - cellular adhesion in treatment of inflammation, perfusion, i metastasis, etc. also for diagnosis.
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The peptide modifies the binding of P-, E- and L-selectin and of integrins, especially intercellular adhesion molecule-1 (ICAM-1). inhibits cellular adhesion. It can be used to treat or prevent
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Misc-difference
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27-SEP-1994
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                                                                    Claim 2; Page 26; 175pp; English.
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                                     Heavner GA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Selectin; integrin; intercellular adhesion molecule; inflammation; coagulation; ischaemia; reperfusion inj
                                                                                                                                                                                                                                                                                                                                                                          Disulfide-bond
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bacterial sepsis; tumour metastisis;
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27-SEP-1994
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CENTOCOR INC
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                                                                                                                                                                                                                                                                                                                                                                                                                          "D-form residue."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tumour;
                                                                                                                                                                     "D-form
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                                                                                                                                                                                                                                                                      "D-form
                                                                                                                                                                                                                                                                                                         "D-form
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 31; DB
Pred. No. 1.9
); Mismatches
                                                                                                                                                                     residue
                                                                                                                                                                                                       residue
                                                                                                                                                                                                                                      residue
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                                                                                                                                                                                                                                                                                                                                                                                          residue."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              leucocyte;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
1.9e+02;
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              imaging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ICAM-1; inhibition;
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RESULT 41
AAR50895
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Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inhibits cellular adhesion. It can be used to treat or prevent inhibits cellular adhesion. It can be used to treat or prevent inhibits cellular adhesion. Ischaemia, reperfusion injury, bacterial sepsis, disseminated intravascular coagulation, adult respiratory distress syndrome, tumour metastisis, rheumatoid arthritis and atherosclerosis. When labelled it can be used to detect defective selectin (or integrin) binding ilgands or high (localised) concentrations of selectin (or integrin) binding cells, especially leucocytes or tumour cells. It can also be used for in vivo imaging. This is a specific example of a degenerate sequence given in the specification. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New cyclic peptide(s) inhibiting selections and integrins - cellular adhesion in treatment of inflammation, perfusion, i
                                                                                                                     Misc-difference
                                                                                                                                              Misc-difference
                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                   Disulfide-bond
Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
27-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The peptide modifies the binding of P-, E- and L-selectin and of integrins, especially intercellular adhesion molecule-1 (ICAM-1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Page 24; 175pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    metastasis, etc. also for diagnosis.
                        WO9405310-A1
                                                                                           Misc-difference
                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                        prevention;
                                                                                                                                                                                                                                                                                                                                                                   disseminated
                                                                                                                                                                                                                                                                                                                                                                                                                                      Cyclic peptide inhibitor of selectins and integrins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR50895 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                 bacterial sepsis; tumour metastisis;
                                                                                                                                                                                                                                                                                                                                                                                                              Selectin;
                                                                                                                                                                                                                                                                                                                                     minated intravascular coagulation; rheumatoid arthritis; respiratory distress syndrome; atherosclerosis; treatment; ntion; detection; tumour; leucocyte; imaging.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                            integrin; intercellular adhesion molecule; ICAM-1; inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                              coagulation;
                                                                                                                                                           /note=
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                                                                                                                                                                                                                                                         "D-form residue."
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                                                                                                                                                                                                                              "D-form residue."
                                                                             "D-form
                                                                                                        "D-form
                                                                                                                                                                                                                                                                                                                                                                                           ischaemia; reperfusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 31; DB 2;
Pred. No. 1.9e+02;
                                                    residue."
                                                                             residue."
                                                                                                        residue
                                                                                                                                 residue."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                             injury;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 - prevent
injury, "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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RESULT 42
AAR50830
ID AAR50
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                integrins, especially intercellular adhesion molecule-1 (ICAM-1). It inhibits cellular adhesion. It can be used to treat or prevent inflammation, coagulation, ischaemia, reperfusion injury, bacterial sepsis, disseminated intravascular coagulation, adult respiratory distress syndrome, tumour metastisis, rheumatoid arthritis and atherosclerosis. When labelled it can be used to detect defective selectin (or integrin) binding ligands or high (localised) concentrations of selectin (or integrin) binding cells, especially leucocytes or tumour cells. It can also be used for in vivo imaging. This is a specific example of a degenerate sequence given in the specification. (Updated on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Page 28; 175pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New cyclic peptide(s) inhibiting selections and integrins - preven cellular adhesion in treatment of inflammation, perfusion, injury,
                            Misc-difference
                                                                            Misc-difference
                                                                                                                               Disulfide-bond
Misc-difference
                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Selectin; integrin; intercellular adhesion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cyclic peptide inhibitor of selectins and integrins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The peptide modifies the binding of P-, E- and L-selectin integrins, especially intercellular adhesion molecule-1 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   metastasis, etc. also for diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1994-100848/12.
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                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                         prevention;
                                                                                                                                                                                                                                                                                                                                                                                                                            disseminated intravascular coagulation, rheumatoid arthritis; adult respiratory distress syndrome; atherosclerosis; treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation; coagulation; ischaemia bacterial sepsis; tumour metastisis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-SEP-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR50830 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2003 to correct PN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-SEP-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4.
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                                                                                                                                                                                                                                                                                                                                                                                                         detection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
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                                                                                                                                                                                       "D-form residue.
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                                                                                                           "D-form
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ellular adhesion molecule; ICAM-1; inhibition; ischaemia; reperfusion injury;
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Pred. No. 1.9e+02;
residue.
                                                     residue.
                                                                                                           residue.
                                                                                                                                                                                                                                                                                                                                                                                                         leucocyte;
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RESULT 43
AAG94724
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New cyclic peptide(s) inhibiting selections and integrins - cellular adhesion in treatment of inflammation, perfusion,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 2; Page 24; 175pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-MAR-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9405310-A1
                          13-DEC-2000; 2000WO-GB004776.
                                                                            14-JUN-2001.
                                                                                                                              WO200142277-A2
                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                             Human;
                                                                                                                                                                                                                                                                                Human
                                                                                                                                                                                                                                                                                                                                     18-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAG94724 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-SEP-1993;
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                                                                                                                                                                                                                                                                          complementary peptide,
                                                                                                                                                                                                                             complementary peptide; ligand; drug discovery; drug design.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-00941653
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                                                                                                                                                                                                                                                                                                                                                                                                                                    peptide; 10
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                                                                                                                                                                                                                                                                          SEQ ID NO: 918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 31; DB 2;
Pred. No. 1.9e+02;
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RESULT 44
AAG94776
ID AAG99
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XX AAG99
XX Humar
XX Humar
XX Humar
XX Homo
XX Ho
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Matches
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                                                                                                                                                                                                                                       A set of peptide ligands consisting of specific complementary peptides proteins encoded by genes of the human genome, useful in an assay for screening and identifying of one or more novel peptides which are drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a set of complementary peptide ligands generated from the human genome. The complementary peptides interact with their relevant target proteins encoded in the human genome. They can be used as reagents in drug discovery and as lead ligands to facilitate drug design and development. The present sequence is a complementary peptide provided and development.
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                                                                                                                                                                                                                                                                                                                                                                                                           Roberts GW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-JUN-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                           Heal JR;
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100.0%; Pr
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID
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The invention relates to a set of complementary peptide ligands generated from the human genome. The complementary peptides interact with their relevant target proteins encoded in the human genome. They can be used as reagents in drug discovery and as lead ligands to facilitate drug design and development. The present sequence is a complementary peptide provided

Page 178;

646pp;

English.

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RESULT 46
AAG86918
ID AAG86
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AC AAG86
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AAG86916
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             AAG86918;
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                                                                                                                                                                                                                                                                                                                              Identifying complementary peptides by analysis of protein and nucleotide sequence databases, useful in drug design.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Saccharomyces cerevisiae; complementary peptide; peptide identification; drug discovery; drug design.
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                                                                                                                                                                                                                                                                                                                                                                                       Roberts GW,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Saccharomyces cerevisiae
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                                 AAG86918 standard; peptide; 10 AA
                                                                                                                                                                                                      Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                         Example 3; Page 287; 488pp; English.
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100.0%; Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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RESULT 47
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Matches 4
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                      17-JUN-2003;
17-JUN-2003;
                                                         17-JUN-2004; 2004WO-EP051158.
                                                                                                                               Unidentified
                                                                                                                                                     diagnosis; cancer; obesity; diabetes; asthma; inflammation; depression;
food; feedstuff; cosmetics; agriculture; animal breeding; VEGF.
                                                                                                                                                                                                                10-MAR-2005
                                                                                                                                                                                                                                        ADV60196;
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10 AA;
                                                                                 23-DEC-2004.
                                                                                                                                                                                                                                                             ADV60196 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Saccharomyces cerevisiae
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                                                                                                        WO2004111636-A2
                                                                                                                                                                                         VEGF peptide SEQ ID NO 7701.
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                      2003EP-00101775.
2003US-0479061P.
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                                                                                                                                                                                                                                                             peptide;
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Pred. No. 1.9e+02;
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reentrrying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the family of protein.
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Bergstrand H,
                                                                    12-APR-1996;
23-SEP-1996;
                                                                                                                       04-APR-1997;
                                                                                                                                                         23-OCT-1997.
                                                                                                                                                                                          WO9739023-A1
                                                                                                                                                                                                                                                              transplant
                                                                                                                                                                                                                                                                                                                                                    27-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                  AAW45003 standard; peptide; 12 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the
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                                                                                                                                                                                                                                                                                                                Immunomodulatory
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                                                                                                                                                                                                                                                            Immunomodulator; immunosuppressant; immunostimulator; treatment;
transplant rejection; autoimmune disease; cancer; infection.
                                   (ASTR ) ASTRA
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 Eriksson T,
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96SE-00003469.
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 Lindvall M,
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Pred. No. 1.9e+02;
   Saernstrand
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunosuppressant can be used to treat transplant rejection or autoimmune disease, e.g. rheumatoid arthritis, systemic lupus crythematosis, Sjogren's syndrome, scleroderma, mixed connective tissue disease, dermatomyositis, polymyositis, Reiter's syndrome, Behcet's disease, type I diabetes, Hashimoto's thyroiditis, Graves' disease, multiple sclerosis, myasthenia gravis, encephalomyelitis, phemphigus vulgaris, vegetans or foliaceus, Senear-Usher syndrome or Brazillan phemphigus. An immunostimulator can be used to treat conditions such as cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleic acids encoding cysteine- or methionine-containing peptide(s) which have immuno:stimulatory or immunosuppressive activity - can be to treat, e.g. cancers, infection, auto:immune disease or transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 12; Page 155; 183pp; English.
                                                                                                                                                                                                                                                                           Synthetic genes
                                                                                                                                                                                                                                                                                                                                                                                 Bergstrand H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9919347-A1
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                                                                                                                                                                                                 Claim 32; Page 52; 104pp;
                                                                                                                                                                                                                                                        methionine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-APR-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunoactive; immunomodulation; immunosuppression; immunostimulation;
immune response; immunoreactive; autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                   (ASTR ) ASTRA
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                                                                                                                                                                                                                                                                                                                                                                                    Lindvall M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 31;
Pred. No.
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                                                                                                                                                                                                                                                                             peptides containing cysteine or
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The present invention describes nucleic acid molecules comprising a coding sequence encoding an immunoreactive peptide and further encoding a protein targeting sequence. The nucleic acid is administered to a patient so that its expression product, an immunoactive peptide, modulates an immune response in a patient. The nucleic acid can also be used to treat cancer, either after surgery to remove a portion of the cancer or after

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conjunction with

the nucleic acid. Cells containing the nucleic acid molecule can

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RESULT 50
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Best Local
              quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosting immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and prognostic or diagnostic disease markers, drug target validation and
                                                                                                                                                                                                                                                                                                  The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants of one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         used for treatment. The immunoactive peptide is immunosuppressive and be used in patients with autoimmune disease. The present sequence represents a specifically claimed immunoactive peptide from the preser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis; cancer; obesity; diabetes; asthma; inflammation; de food; feedstuff; cosmetics; agriculture; animal breeding; VEGF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-JUN-2003; 2003EP-00101775.
17-JUN-2003; 2003US-0479061P.
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profiling, pathway
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cc corresponds with a family of proteins where each of the members of the cpeptide combo is derived from a unique protein from the family. The cc peptide combo is useful for quantifying specific known splice variants of cc one or more particular proteins in a sample, for diagnosing complex cc genetic diseases such as cancer, obesity, diabetes, asthma and cc inflammation, neuropsychiatric discorders such as depression, for cc quantifying one to several hundreds of protein disease markers cc simultaneously leading to a more accurate diagnostic sub-classification, cc for determining the extent of protein modification in a particular sample cf for determining the extent of protein modification in a particular sample cc of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases co rneurological diseases, as biomarkers preclinical drug development, development, clinical drug development, guidance marketed drugs, cc rosicology, clinical drug development, guidance marketed drugs, cc prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway cc analysis, answering basic disease biology questions, and in the fields of cool and feed, cosmetics, agriculture and animal breeding. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 51
ADV60192
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Best Local S
Matches 4
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food and f
                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-JUN-2003; 2003EP-00101775
17-JUN-2003; 2003US-0479061P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADV60192 standard; peptide; 13
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                                                                                                                                                                                                                                                                                                                                                                                           Example; SEQ ID NO 7697; 265pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2005-057893/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kas K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADV60192;
                                                                                                                                                                                                                                                                                                                                                           The invention relates to
                                                                                                                                                                                                                                                                                                                                                                                                                              family of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-JUN-2004; 2004WO-EP051158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VEGF peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYGE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (VIBV-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                psis; cancer; obesity; diabetes; asthma; inflammation; de
feedstuff; cosmetics; agriculture; animal breeding; VEGF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UNIV GENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VIB VZW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       feed, cosmetics,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  answering basic disease biology questions, and in the fields feed, cosmetics, agriculture and animal breeding. The present represents a peptide from a VEGF peptide combo.
                                                                                                                                                                                                                                                                                                                                                                                                                              protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7697
                                                                                                                                                                                                                                                                                                                                                           a method of identifying
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                                                                                                                                                                                                                                                                                                                                        of
                                                                                                                                                                                                                                                                                                                                                           peptide combo which
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RESULT 52
ADV60193
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                       quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present
                                                                                                                                                                                                                                                     The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diagnosis; cancer; obesity; diabetes; asthma; inflammation; depression; food; feedstuff; cosmetics; agriculture; animal breeding; VEGF.
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17-JUN-2003; 2003US-0479061P.
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                                                                                                                                                                                                                                                                                                                                                                                                Example; SEQ ID NO 7698;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2005-057893/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                 family of protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vandekerckhove
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                                                                                                                                                                                                                                                                                                                                                                                               265pp; English
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Pred. No. 2.3e+02;
                  from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>-</u>
              a VEGF peptide
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RESULT 53
ADW28625
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          CC target proteins. The method comprises: (i) contacting an antibody-coated CC solid surface with a fusion protein, where the antibodies bind CC specifically to the fusion protein and the fusion protein comprises at CC least a portion of the target protein linked to a carrier protein, and CC (ii) conducting an assay to determine the presence of the carrier CC protein, where the presence of a carrier protein indicates the presence CC of an antibody to the target protein. The target protein may be an CC isoform of a protein that is associated with disease, e.g. Vascular CC Endothelial Growth Factor (VEGF) isoforms VEGF165 (ADW28643) and VEGF121 CC (ADW28645), which are associated with prostate cancer and Her2 CC (ADW28645), which is associated with prostate cancer and Her2 CC (ADW28640), which is associated with prostate cancer and Her2 CC (ADW28640) binds to a target protein, specifically at least one monoclonal CC antibody binds to a target protein, specifically at least one monoclonal CC disease; isolating (M2) monoclonal antibodies, where each monoclonal antibody that binds to an isoform of a protein that is associated with a CC disease; isolating (M3) an antibody binding specifically to a target protein from several antibodies that are associated with the nucleic CC acid(s) encoding the antibodies that are associated with the concerning CC antibods identified by (M1) are useful for tracting or preventing CC diseases in which the presence of an antibody to a particular molecule is beneficial. The antibodies are useful for tracting according to a concerning constitution of a concerning concerning constitution of a concerning concerning conc
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying antibody to protein, involves contacting solid surface coated with antibodies with fusion protein having portion of target protein and carrier protein and conducting assay to determine presence of carrier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytostatic; Antirheumatic; Antidiabetic; Cardiant; Nootropic; Neuroprotective; antibody identification; antibody production; antibody therapy; cancer; vascular endothelial cell growth factor; VEGF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 32; SEQ ID NO 2; 55pp; English.
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to particular cells e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for identifying antibodies for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein.
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cancer cell.
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RESULT 54
ADV54876
ID ADV54
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The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants or one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   epitopes on target protein which are used for preparing DNA vaccines having nucleotide sequences encoding epitope of a disease associated protein isoforms used for prevention or treatment of diseases such as cancer, rheumatoid arthritis, diabetes, acute myeloid leukemia (CLL), chronic lymphocytic leukemia (CLL), ovarian cancer, prostate cancer, cardiovascular disease, Alzheimer's disease, etc. The antibodies are useful for detecting antigens e.g. in specific cells, tissue or bodily fluids such as serum, or for affinity purification of antigen from recombinant cell culture or natural sources. The present sequence is a VEGF isoform epitope, which can be used to produce a VEGF isoform fusion
                                                                                                                                                                                                                                                                                 Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the family of protein.
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17-JUN-2003; 2003US-0479061P
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                       The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants of one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
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17-JUN-2003; 2003US-0479061P
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Pred. No. 2.8e+02;
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The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants of one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development,
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food; feedstuff; compared
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cosmetics; agriculture; anima
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          combo which corresponds with a family of protein a variety of diseases, drug development or in generating peptides by applying a digest on the
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Pred. No.
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ches 0;
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RESULT 57
AAB23038
ID AAB23
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cardiovascular disease; angina pectoris; arteriosclerosis; heart failure; brain disorder; brain abscess; meningitis; Alzheimer's disease; sarcoma; cytostatic; carcinoma; cardiant; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAE23038 standard; peptide; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer; lung; breast;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            thioredoxin,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             head; neck;
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The invention relates to human thioredoxin nucleic acid and polypeptide molecules, designated 22108 and 47916. The compound that modulates the activity or expression of 22108 and 47916 nucleic acid is useful for treating or preventing a disorder characterised by aberrant activity of 22108 and 47916-expressing cell, specifically for reducing or inhibiting the aberrant activity of the 22108 and 47916 nucleic acid and polypeptide are useful for diagnosing, preventing or treating cancer in a subject (e.g. carcinoma, sarcoma, metastatic or haematopoietic disorders (e.g.leukaemia), or cancers of the lung, breast, thyroid, head neck, prostate or genito-urinary tract), cardiovascular diseases (e.g. angina pectoris, arteriosclerosis or heart failure) or brain disorders (e.g. brain abscess, meningitis, Alzheimer's diseases). The thioredoxin DNA is also useful in gene therapy. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New human thioredoxin nucleic acid and polypeptide molecules, designated 22108 and 47916, useful for diagnosing, preventing or treating cancer (e.g. carcinoma), cardiovascular diseases (e.g. heart failure) or brain
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WPI; Bandaru R, 25-SEP-2000;

2000US-0235049P

25-SEP-2001; 2001WO-US029967

(MILL-) MILLENIUM PHARM INC

Kapeller-Libermann

04-APR-2002 WO200226803-A2

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ADV60188
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     quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present sequence represents a peptide from a VEGF peptide combo.
                                                                                                                                                                                                                                                                corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the family of protein.
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17-JUN-2003; 2003US-0479061P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diagnosis; cancer; obesity; diabetes; asthma; inflammation; depression; food; feedstuff; cosmetics; agriculture; animal breeding; VEGF.
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                                                                                                                                                                                                                                                                                                                                                           The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the
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                                                                                                                                                                                                                                                                                                                                                                                                              Example; SEQ ID NO 7693; 265pp; English.
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Pred. No. 3.1e+02;
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Best Local S
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                                       quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present
                                                                                                                                                                                                                                                             The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants of one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and inflammation, neuropsychiatric disorders such as depression, for
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17-JUN-2003; 2003US-0479061P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       VEGF peptide SEQ ID NO 7694.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADV60189 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-DEC-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (VIBV-) VIB VZW.
                                                                                                                                                                                                                                                                                                                                                                                                                                  family of protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         feedstuff; cosmetics;
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                             represents a peptide from a VEGF peptide combo
                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 7694; 265pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CGPC 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diabetes; asthma; inflammation; depression; agriculture; animal breeding; VEGF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21
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Pred. No. 3.2e+02;
); Mismatches 0;
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Best Local Similarity
Matches 4; Conser
                                                                                                                  invention. Human secreted proteins have activities based on the tissues and cells the genes are expressed in. Example of activities include: immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; notropic; cytostatic; cardiant; vasotropic; cerebroprotective; notropic; operative; notropic; cytostatic; antibacterial; virucide; fingicide; and ophthalmological. The polynucleotides and polypeptides can be are used to prevent, treat or ameliorate a medical condition in e.g. humans, mice, rabbits, goats, horses, cate, dogs, chickens or sheep. They are also used in diagnosing a pathological condition or susceptibility to a pathological condition or susceptibility to a pathological condition. Disorders which are diagnosed or treated include pathological condition. Disorders which are diagnosed or treated include autoimmune diseases, hyperproliferative disorders, cardiovascular disorders, infections caused by bacteria, viruses and fungi and ocular disorders. The polypeptides can also be used to aid wound healing and disorders. The polypeptides can also be used to aid wound healing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; ophthalmological; gene therapy; autoimmune disease; infection; hyperproliferative disorder; cardiovascular disorder; angiogenesis; cerebrovascular disorder; nervous system disorder; ocular disorder; wound healing; skin aging; food additive; preservative.
                                disorders. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sumburn, maintain organs before transplantation, for supporting cell culture oprimary tissues, to regenerate tissues and in chemotaxis. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human secreted proteins given in AAB38119 to AAB38165. AAB38166 to AAB38201 represent human secreted polypeptide sequences and proteins homologous to them, which are given in the exemplification of the pro-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The polynucleotide sequences given in AAC69399 to AAC69445 encode human secreted proteins given in AAB38119 to AAB38165. AAB38166 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acids encoding human secreted proteins, used to prevent, ameliorate, or diagnose conditions such as autoimmune disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-1999;
22-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB38160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-611713/58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-MAR-2000; 2000WO-US007526
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ruben SM,
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99US-0171550P
   also
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   Corresponds with a family of proteins where each of the members of the CC corresponds with a family of proteins where each of the members of the CC peptide combo is derived from a unique protein from the family. The CC peptide combo is useful for quantifying specific known splice variants of CC one or more particular proteins in a sample, for diagnosing complex CC quantifying one to several hundreds of protein disease markers CC quantifying one to several hundreds of protein disease markers CC simultaneously leading to a more accurate diagnostic sub-classification, for constraining the extent of protein modification in a particular sample CC of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels CC diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, CC coxicology, clinical drug development, guidance marketed drugs, CC propositic or diagnostic disease markers, drug target validation and CC selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
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analysis, a...
and feed,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G protein coupled
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-JUN-2004; 2004WO-EP051158
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17-JUN-2003; 2003US-0479061P
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RESULT 62
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ID ADV56
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 The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the combotiants of the combo is useful for quantifying specific known splice variants of come or more particular proteins in a sample, for diagnosing complex complex comparation, neuropsychiatric disorders such as descript, diabetes, asthma and committying one to several hundreds of protein disease markers committed in the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for grenatal testing to detect the presence of a congenital disease or for quantitating protein levels corneurological diseases, as biomarkers preclinical drug development, cornection, monitoring protein models, biomarkers related with cornection, monitoring protein splicing, drug tearget validation and selection, monitoring protein splicing, drug lead profiling, pathway cannot represent a correction complex resents a mentide freed, cosmetice, agriculture and animal breeding. The present
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Best Local :
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17-JUN-2003; 2003US-0479061P.
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stuff; cosmetics;
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Pred. No.
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           The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The composite combo is useful for quantifying specific known splice variants of the genetic diseases such as cancer, obesity, diabetes, asthma and complex compl
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Best Local Similarity
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17-JUN-2003; 2003US-0479061P
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) UNIV GENT.
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Pred. No. 3.6e+02;
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                                          The invention relates to a method of identifying a peptide combo which corresponds with a family of proteains where each of the members of the peptide combo is derived from a unique protein from the family. The composition is useful for quantifying specific known splice variants of come or more particular proteins in a sample, for diagnosing complex competite diseases such as cancer, obesity, diabetes, asthma and continuously leading to a more accurate diagnostic sub-classification, for comparatifying one to several hundreds of protein disease markers comparatifying one to several hundreds of protein as depression, for comparatifying one to several hundreds of protein a particular sample of retermining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels cor neurological diseases, as biomarkers preclinical drug development, contical drug development, guidance marketed drugs, clinical drug development, guidance marketed drugs, answering basic disease markers, drug target validation and constitute, agriculture and animal breeding. The present contical constitutes, agriculture and animal breeding. The present contical constitutes, agriculture and animal breeding.
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17-JUN-2003; 2003US-0479061P.
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4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                               of protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vandekerckhove J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    UNIV GENT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Krols L;
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                              G-protein
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RESULT 65
ADV55148
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CC genetic diseases such as cancer, obesity, diabetes, asthma and confirmation, neuropsychiatric disorders such as depression, for cquantifying one to several hundreds of protein disease markers cc simultaneously leading to a more accurate diagnostic sub-classification, cf proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels cd diagnostic of a chromosomal abnormality, for diagnosing immune diseases cor neurological diseases, as biomarkers preclinical drug development, cd development of improved animal models, biomarkers related with coxicology, clinical drug development, guidance marketed drugs, compensatic or diagnostic disease markers, drug terget validation and compensation, monitoring protein splicing, drug lead profiling, pathway canalysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present compensations are represents a peptide from a G-protein coupled receptor peptide
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Best Local S
Matches 4
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                                   sequence combo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kas K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G protein coupled receptor peptide SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-JUN-2003; 2003EP-00101775.
17-JUN-2003; 2003US-0479061P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis; cancer; obesity; diabetes; asthma; inflammation; depression;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         family of protein.
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100.0%; Pred. No. 3.6e+02;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Krols L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24
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Sequence 24 AA

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RESULT 66
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                                                                                                                                     Sequences AAB37984-B38019 represent the amino acid sequences of 27 human compared by the genes AAC69084-C69119. The genes and proteins are useful for preventing, ameliorating or treating medical conditions, e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the college of a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the college ovarian cancer, and other cancers of the adrenal gland, bone, bone immune discreter, gastrointestinal tract, liver, lung, or urogenital; (b) immune discreters e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, allergies, autoimmune haemolytic cardiovascular disorders such as mycocardial ischaemias; (d) wound healing (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral, bacterial, fungal and parasitic
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Best Local Similarity
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                                                Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                 New nucleic acid molecules encoding 27 human secreted proteins for diagnosing, preventing, treating or ameliorating medical conditions and used as food additives or preservatives.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer; vulnerary, antionvulsant; antibacterial; antifungal; antiparastic; cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cardiant; gene therapy; cancer; immune disorder; cardiova neurological disease; infection; human; secreted protein.
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                                                                                              Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
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   2 CGPC 5
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s G, Lafleur DW,
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                                                                                                                                                                                                                                                                                                                                                                                                     Page 20;
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Moore PA,
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Pred. No. 3.6e+02;
                                                Score 31; DB 3; 1
Pred. No. 3.8e+02;
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                                 Mismatches
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Olsen
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HS, Young
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Mismatches

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3.8e+02; DB 9;

Length 25; Indels

100.0%; Score 31; 100.0%; Pred. No.

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                                                                                          quantifying one to several hundreds of protein disease markers simultaneously leading to a more accurate diagnostic sub-classification, for determining the extent of protein modification in a particular sample of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, development of improved animal models, biomarkers related with toxicology, clinical drug development, guidance marketed drugs, prognostic or diagnostic disease markers, drug target validation and selection, monitoring protein splicing, drug lead profiling, pathway analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present sequence represents a peptide from a G-protein coupled receptor peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The peptide combo is useful for quantifying specific known splice variants one or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obseity, diabetes, asthma and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genetic diseases such as cancer, obesity, diabetes, asthma an inflammation, neuropsychiatric disorders such as depression,
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Sequence
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17-JUN-2003; 2003US-0479061P.
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25 AA;
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                                                                                                                                                                                               The invention relates to a method of identifying a peptide combo which corresponds with a family of proteins where each of the members of the peptide combo is derived from a unique protein from the family. The composition is useful for quantifying specific known splice variants of the peptide combo is useful for quantifying specific known splice variants of come or more particular proteins in a sample, for diagnosing complex genetic diseases such as cancer, obesity, diabetes, asthma and complex complex complex genetic diseases such as cancer, obesity, diabetes, asthma and comparison one to several hundreds of protein disease markers comparison one to several hundreds of protein disease markers comparison of the extent of protein modification in a particular sample for determining the extent of protein modification in a particular sample cof proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels cor neurological diseases, as biomarkers preclinical drug development, constituted in the fields of toxicology, clinical drug development, guidance marketed drugs, answering basic disease biology questions, and in the fields of cond and feed, cosmetics, agriculture and animal breeding. The present
                                                                                 Query Match
Best Local Similarity
                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-JUN-2003; 2003EP-00101775
17-JUN-2003; 2003US-0479061P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnosis; cancer;
food; feedstuff; co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G protein coupled receptor peptide SEQ ID NO 2537.
                                                                                                                                   Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example;
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                                                                                                                                                                    sequence represents a peptide combo.
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æ
                                 2 CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 2537; 265pp; English.
CGPC
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                                                                 Conservative
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                                                                                 100.0%;
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                                                                 0
                                                                                 Score 31; DB 9;
Pred. No. 3.8e+02;
                                                                                                                                                                                    from a G-protein coupled receptor peptide
                                                                 Mismatches
                                                                                                 Length 25;
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RESULT 69
ADV54928
                                                                                                                                                                                   The invention relates to a method of identifying a peptide combo which CC corresponds with a family of proteins where each of the members of the Peptide combo is derived from a unique protein from the family. The CC peptide combo is useful for quantifying specific known splice variants of one or more particular proteins in a sample, for diagnosing complex CC genetic diseases such as cancer, obesity, diabetes, asthma and CC inflammation, neuropsychiatric disorders such as depression, for CC quantifying one to several hundreds of protein disease markers CC simultaneously leading to a more accurate diagnostic sub-classification, CC for determining the extent of protein indistance markers CC of proteins, for tissue-typing analysis, for prenatal testing to detect the presence of a congenital disease or for quantitating protein levels CC diagnostic of a chromosomal abnormality, for diagnosing immune diseases or neurological diseases, as biomarkers preclinical drug development, CC development of improved animal models, biomarkers related with coxicology, clinical drug development, guidance marketed drugs, CC genbo, answering basic disease markers, drug target validation and CC selection, monitoring protein splicing, drug lead profiling, pathway CC analysis, answering basic disease biology questions, and in the fields of food and feed, cosmetics, agriculture and animal breeding. The present crowders appetide from a c-protein coupled receptor peptide cc combo.
Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying a peptide combo which corresponds with a family of proteins, useful for diagnosing a variety of diseases, drug development or in agriculture, comprises generating peptides by applying a digest on the
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17-JUN-2003; 2003US-0479061P.
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                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (VIBV-) VIB VZW.
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   100.0%;
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   Score 31; DB 9;
Pred. No. 3.8e+02;
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                                   Length 25;
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CGPC 11

Matches

4. Similarity

Conservative

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Mismatches

Indels

0

Gaps

0

2 CGPC 5

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RESULT 70
ADC89744
ID ADC8974
ID 
                                                                                                                                                                                                    The invention relates to a cyclic VEGF (vascular endothelial growth CC factor) peptide (ADC89740) with NP-1 (neuropilin-1) antagonist activity. CC The peptide corresponds to residues 138-165 of the VEGF-165 splice CC variant, but contains 2 disulphide bonds. It acts as a competitive CC (including VEGF-mediated tumour angiogenesis), neurodegeneration CC (involving interaction with semaphorin 3) and in immunosuppression. NP-1 (is a non-tyrosine kinase receptor for the VEGF-165 splice variant. CC Additionally, VEGF is known to promote the survival of tumour cells that CC express NP-1, including some breast carcinoma cells. The cyclic VEGF CC peptide of the invention can be used to stimulate nerve growth and CC repair, to treat neurodegeneration, and for anti-cancer therapy. It can C also be used for treating other angiogenic conditions such as eye CC diseases and rheumatoid arthritis, and as an immunosuppressant for use in CC transplantation. The present sequence represents a cyclic VEGF-165 CC derived peptide which was used in a study of VEGF-165 binding to NP-1 (CC expressing cells. The present sequence had no inhibitory effect on the CC binding of VEGF-165 to NP-1, unlike the cyclic peptide of the invention CC (ADC89740).
Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                              Sequence 28 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New cyclic peptide derived from vascular endothelial growth factor, useful for treating e.g. neurodegeneration and cancer, antagonizes neuropilin-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-804028/75.
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Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-MAR-2003; 2003WO-GB001375
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Disulfide-bond
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cyclic VEGF-165-derived peptide (residues 111-138), 3286.
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              Conservative
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          100.0%; Score 31; DB 7; 1 100.0%; Pred. No. 4.1e+02; /ative 0; Mismatches 0;
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A;Description: Purification and Sequence deter
A;Reference number: A59356
A;Accession: A59356
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-35 <XUC-
A;Cross-references: UNIPARC:UPI00001766C6
C;Superfamily: scorpion neurotoxin
C;Keywords: venom
F;1-35/Product: neurotoxin BmK 41-2 #status ex
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C;Species: Androctonus mauretanicus mauretanicus
C;Date: 15-Oct-1982 #sequence_revision 15-Oct-1982 #text_change 09-Jul-2004
C;Accession: A01758
R;Rosso, J.P.; Rochat, H.
Toxicon 23, 113-125, 1985
A;Fitle: Characterization of ten proteins from the venom of the Moroccan scorpion Androc A;Reference number: A94318; MUID:85193276; PMID:3992595
A;Accession: A01758
A;Accession: A01758
A;Residues: 1-35 - ROS>
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C;Superfamily: scorpion neurotoxin
C;Keywords: neurotoxin; venom
F;1-18,4-25,15-30,19-32/Disulfide bonds: #status predicted
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;Date: 31-Dec-2001 #sequence_revision 31-Dec-2001 #text;Accession: A59356
;1-35/Product: neurotoxin BmK 41-2 #status experimental;1-18,4-25,15-30,19-32/Disulfide bonds: #status predicte
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testis-specific protein (clone mst(3)gl-9) - fruit (;Species: Drosophila melanogaster C;Date: 30-Sep-1990 #sequence_revision 30-Sep-1990 C;Accession: S00340 R;Kuhn, R.; Schaefer, U.; Schaefer, M. EMBO J. 7, 447-454, 1988 A;Fitle: Cis-acting regions sufficient for spermato A;Fitle: Cis-acting regions sufficient for spermato A;Fitle: Cis-acting regions MUID:88211557; PMID:283
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C;Genetics:
A;Gene: Mst84Dc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cestis specific protein Mst84Dc - fruit fly (Drosophila melanogaster)
C;Species: Drosophila melanogaster
C;Date: 26-Uul-1996 #sequence_revision 26-Uul-1996 #text_change 09-Uul-2004
C;Accession: S25774; C56565
R;Kuhn, R.; Kuhn, C.; Boersch, D.; Glaetzer, K.H.; Schaefer, U.; Schaefer, Mech. Dev. 35, 143-151, 1991
A;Title: A cluster of four genes selectively expressed in the male germ line A;Reference number: A56565; MUID:92102953; PMID:1684716
A;Accession: S25774
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C;Superfamily: fruit fly
C;Keywords: spermatogenes
                                                                                                                                       A;Cross-references: FlyBase:FBgn0002862
C;Superfamily: fruit fly testis-specific
C;Keywords: sex-specific protein; testis
                                                                                                                                                                                                                                                      A; Molecule type: DNA
A; Residues: 1-56 < KUH>
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A; Residues: 1-55 < KUH>
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Pred. No. 81;
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rubredoxin NMB0993 [imported] - Neisseria meningitidis (strain MC58 serogroup C;Species: Neisseria meningitidis G;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text chance no Total C;Accession: H81131, Balloo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPARC:UPI00000C4E89; GB:AL162755; GB:AL157959; NID:g7379742; PIDN A;Experimental source: serogroup A, strain Z2491
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Title: Complete DNA sequence of a serogroup A strain of Neisseria menigitidis Z2491.
A;Reference number: A81775; MUID:20222556; PMID:10761919
A;Accession: B81888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPROT:Q9JQN3; UNIPARC:UPI00000C4E89; GB:AE002450; GB:AE002098; NII A;Experimental source: serogroup B, strain MC58 R;Parkhill, J.; Achtman, M.; James, K.D.; Bentley, S.D.; Churcher, C.; Klee, S.R.; Morel; Holroyd, S.; Jagels, K.; Leather, S.; Moule, S.; Mungall, K.; Quail, M.A.; Rajandream, Nature 404, 502-506, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Accession: H81133; B81888

R;Tettcelin, H.; Saunders, N.J.; Heidelberg, J.; Jeffries, A.C.; Nelson, K.E.; Eiser Hickey, E.K.; Haft, D.H.; Salzberg, S.L.; White, O.; Fleischmann, R.D.; Dougherty, ri, H.; Qin, H.; Vamathevan, J.; Gill, J.; Scarlato, V.; Masignani, V.; Pizza, M. Science 287, 1809-1815, 2000

A;Authors: Grandi, G.; Sun, L.; Smith, H.O.; Fraser, C.M.; Moxon, E.R.; Rappuoli, R.A;Title: Complete genome sequence of Neisseria meningitidis serogroup B strain MC58 A;Reference number: A81000; MUID:20175755; PMID:10710307

A;Accession: H81133
                    RESULT
S25772
                                                                                                                                                                                                                                                                                                                                                                                    J. Mol. Endocrinol. 4, 143-150, 1990
A;Title: Partial cDNA sequence for the donkey chorionic A;Reference number: 815676; MUID:90262634; PMID:2344391
A;Accession: $15676
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C;Date: 21
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C;Superfamily: rubredoxi
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A; Residues: 1-56 < TET>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Accession: S15676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chorionic gonadotropin beta chain -
testis-specific protein Mst84Da -
                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                           Cross-references: UNIPROT:P19794; UNIPARC:UPI00001765CA; EMBL:X53669;Superfamily: pituitary glycoprotein hormone beta chain
                                                                                                                                                                                                                                                                                                                        ;Molecule type: mRNA
;Residues: 1-62 <LEI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Species: Equus asinus (donkey)
Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
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Residues: 1-56 <PAR>
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fruit fly (Drosophila melanogaster)
                                                                                                                                                                                                               Score 31;
Pred. No.
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Pred. No.
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                                                                                                                                                                                             Mismatches
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erty, B.A.
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A;Accession: A60136
A;Molecule type: mRN
A;Residues: 1-73 <WJ
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Mech. Dev. 35, 143-151,
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A; Residues: 1-63 < KUH>
                                                                                                                                                C; Accession: A60136
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A; Residues: 1-68 < KUH>
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                                                                                                                                                                                                                                                           RESULT 9
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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C;Species: Drosophila melanogaster
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 09-Jul-2004
C;Accession: S25772; A56565
R;Kuhn, R.; Kuhn, C.; Boersch, D.; Glaetzer, K.H.; Schaefer, U.; Schaefer, M.
Mech. Dev. 35, 143-151, 1991
A;Title: A cluster of four genes selectively expressed in the male germ line of Drosoph
A;Reference number: A56565; MUID:92102953; PMID:1684716
Biochim. Biophys. Acta 824, 201-208, 1985
A;Title: Isolation and characterisation of keratin mRNA
A;Reference number: A60136; MUID:85122780; PMID:2578818
                                                                                                                                        keratin, scale (clone CSK9) - chicken (fragment)
C;Species: Gallus gallus (chicken)
C;Date: 30-Jan-1993 #sequence_revision 12-Mar-19
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C;Superfamily: fruit fly testis-specific protein
C;Keywords: spermatogenesis; tandem repeat
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A;Accession: S25775
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C;Species: Drosophila melanogaster
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change
C;Accession: $25775; D56565
                                                                                      R; Wilton, S.D.; Crocker, L.A.; Rogers,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPROT:Q01645; UNIPARC:UPI000012EAE7; EMBL:X67703; NID:g11072; A;Note: sequence extracted from NCBI backbone (NCBIN:74217, NCBIP:74223)
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ermatogenesis; tandem repeat
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3-151, 1991
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100.0%; Pred. No. 95;
tive 0; Mismatches
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    fruit fly (Drosophila melanogaster)

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C;Species: Drosophila melanogaster
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 09-Jul-2004
C;Accession: S25773; B56555
R;Kuhn, R.; Kuhn, C.; Boersch, D.; Glaetzer, K.H.; Schaefer, U.; Schaefer, M.
Mech. Dev. 35, 143-151, 1991
Mech. Dev. 35, 143-151, 1991
A;Title: A cluster of four genes selectively expressed in the male germ line of A;Reference number: A56565; MUID:92102953; PMID:1684716
A;Reference number: A56565; MUID:92102953; PMID:1684716
A;Accession: S25773
A;Molecule type: DNA
A;Residues: 1-74 <KUH>
                                                                                                                                                                                                                                                                                                              C;Species: Rattus norvegicus (Norway rat)
C;Date: 29-May-1998 #sequence_revision 29-May-1998 #text
C;Accession: I65235
R;Zhang, F.P.; Rannikko, A.; Huhtaniemi, I.
Biochem. Biophys. Res. Commun. 210, 858-865, 1995
Biochem. Biophys. Res. Commun. 210, 858-865, 1995
A;Title: Isolation and characterization of testis-specif
A;Reference number: I52320; MUID:95283549; PMID:7763258
A;Accession: I65235
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C;Superfamily: feather keratin
C;Keywords: tandem repeats
F;1-49/Region: 13-residue repeats
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165235
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A;Map position: 3
C;Superfamily: fruit fly testis-specific protein C;Keywords: spermatogenesis; tandem repeat
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C;Species: Drosophila melanogaster
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 09-Ju
                                                                S
                                                                                                                                                                                                                                                          A;Status: preliminary; translated A;Molecule type: mRNA A;Residues: 1-80 <RES>
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C;Species: Rattus norvegicus (Norway rat)
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Best Local (
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Best Local
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Pred. No.
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                                                                                                         Score 31; DB 2;
Pred. No. 1.1e+02;
Mismatches 0;
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Pred. No.
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A;Experimental source: calamus
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thioredoxin 2 - slime mold (Dictyostelium discoideum) (fragment) C;Species: Dictyostelium discoideum C;Species: Dictyostelium discoideum C;Date: 28-Oct-1994 #sequence_revision 28-Oct-1994 #text_change C;Accession: B46264 R;Wetterauer, B.; Jacquot, J.P.; Veron, M. J. Biol. Chem. 267, 9895-9904, 1992 A;Title: Thioredoxins from Dictyostelium discoideum are a develo; A;Reference number: A46264; MUID:92250653; PMID:1577820 A;Accession: B46264
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E84409
                                                                                                                                                                                                                       KRGLBS

Keratin, feather - silver gull

keratin, feather - silver gull

C;Species: Larus novaehollandiae (silver gull)

C;Species: 1204 #secuence revision 24-Apr-1984
                                                                                                                                                                                                                                                                                               RESULT 14
KRGLBS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Leithauser, B.; Keller, K.; Cruz, R.; Danson, M.J.; Hough, Jung, K.H.; Alam, M.; Freitas, T.

Proc. Natl. Acad. Sci. U.S.A. 97, 12176-12181, 2000

A;Authors: Hou, S.; Daniels, C.J.; Dennis, P.P.; Omer, A.D.;

A;Title: Genome sequence of Halobacterium species NRC-1.

A;Reference number: A84160; MUID:20504483; PMID:11016950
                                                                                      C;Accession: A02850

R;O'Donnell, I.J.; Inglis, A.S.

R;Accession: A02850; MUID:75054009; PMID:4429491

A;Accession: A02850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Gene: trxA2
C;Superfamily: thioredoxin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Halobacterium sp. NRC-1
C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 05-Oct-2004
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A; Residues: 1-88 <WET>
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                         A;Residues: 1-98 <OI
A;Cross-references:
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A; Residues: 1-89 <STO>
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                                              A;Molecule type: protein
A;Residues: 1-98 <ODO>
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UNIPROT: P02451; UNIPARC: UPI000012E06F
be: calamus of wing feathers from a sir
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Pred. No. 1.2
0; Mismatches
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                                                                                                                                       gull (Larus novae-holland
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).G.; Jable
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R;Shirai, M.; Hirakawa, H.; Kimoto, M.; Tabuchi, M.; Kishi, F.; Ouchi, K.; Shiba, Nucleic Acids Res. 28, 2311-2314, 2000
A;Title: Comparison of whole genome sequences of chlamydia pneumoniae J138.
A;Reference number: A86491; MUID:20330349; PMID:10871362
A;Accession: H86572
                                                                                                                                                                                                                                                                                                                      thioredoxin [imported] - Chlamydophila pneumoniae (strain J138) C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change C;Accession: H86572
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                                                                                                                                    A;Cross-references: UNIPROT:Q9Z7P5; UNIPARC:UPI000004461A; A;Experimental source: strain J138
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A;Experimental source: strain
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Science 270, 397-403, 1995
A;Tille: The minimal gene complement of Mycoplasma genitalium.
A;Reference number: A64200; MUID:96026346; PMID:7569993
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A; Residues: 1-102 <STO>
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Accession: G64213
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                  100.0%;
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Pred. No.
                  Score 31; DB
Pred. No. 1.30
    Mismatches
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.3e+02;
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                                                                                                                                                      GB:BA000008; NID:g8979031;
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thioredoxin CP0088 [imported] - Chlamydophila pneumoniae (strains CWL029 and AR39) C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C:Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 05-Oct-2004 C;Accession: D72052; E81614
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B71503
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Science 282, 754-759, 1998
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A; Residues: 1-102 < ARN>
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A; Residues: 1-102 <HIM>
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Best Local S
Matches 4
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A;Gene: trxA
C;Superfamily: thioredoxin; thioredoxin homology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R; Himmelreich, R.; Hilbert, H.; Plagens, H.; Pirkl, E.; Li, B.C.; Herrmann, R. Nucleic Acids Res. 24, 4420-4449, 1996
A;Title: Complete sequence analysis of the genome of the bacterium Mycoplasma A;Reference number: S73327; MUID:97105885; PMID:8948633
A;Accession: S73896
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C;Superfamily: thioredoxin; thioredoxin homology
F;9-90/Domain: thioredoxin homology <TXN>
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A;Variety: ATCC 29342
C;Date: 27-Feb-1997 #sequence_rev
C;Accession: $73896
                                                                                                                                                                      A;Cross-references: UNIPROT:O84544; UNIPARC:UPI0000136E75; GB:AE001324; GB:AE001273; NID
A;Experimental source: serotype D, strain UW-3/Cx
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                                                                                                                                                                                                                                                                                                                   A; Reference number: A71570;
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100.0%; Score 31; Dilarity 100.0%; Pred. No. 1.:
Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                   of an obligate intracellular pathogen 70; MUID:99000809; PMID:9784136
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Pred. No. 1.3e+02;
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                         1.3e+02
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thioredoxin II - yeast (Saccharomyces N;Alternate names: protein L1933; protein C;Species: Saccharomyces cerevisiae C;Dete: 31-Dec-1991 #sequence_revision C;Accession: S15048; A39847; S15360; % R;Gan, Z.R.

J. Biol. Chem. 266, 1692-1696, 1991
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C;Superf
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                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPROT:Q9PJK3; UNIPARC:UPI0000057A90; GB:AE002349; GB:AE002160; A;Experimental source: strain Nigg (MoPn)
C;Genetics:
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C;Species: Chlamydia muridarum, Chlamydia trachomatis MoPn
C;Date: 31-Max-2000 #sequence_revision 31-Max-2000 #text_change 05-Oct-2004
C;Accession: C81660
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A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39.
A;Reference number: A81500; MUID:20150255; PMID:10684935
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A;Status: preliminar
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A; Residues: 1-102 < REA>
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                                         ;Date: 31-Dec-1991 #sequence_revision;Accession: S15048; A39847; S15360; S6
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Best Local
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Best Local
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                                                                                                                              yeast (Saccharomyces cerevisiae)
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Pred. No. 1.
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Pred. No. 1.36
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                                             S64870; B38669
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T39085
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A; Residues: 1-103 < MUL>
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J. Biol. Chem. 2
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A; Residues: 1-103 < KOE>
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                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-103 <GEN>
                                                                                                                                                                                                                                                                           A; Accession: T39085
                                                                                                                                                                                                                                                                                              A; Reference number: Z21826
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                          A;Molecule type: mRNA
                                                                   A; Accession:
                                                                                    A; Reference number: Z23075
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A;Reference number: A38669; MUID:91107668; PMID:1988444
A;Reference number: A38669; MUID:91107668; PMID:1988444
A;Accession: S15048
A;Accession: LONA
A;Residues: 1-103 <GAN>
A;Residues: 1-103 <GAN>
A;Cross-references: UNIPROT:P22217; UNIPARC:UPI000004F943; EMBL:M59169; NID:g173027; PII
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R;Koetter, P.; Rose, M.; Entian, K.D.
submitted to the Protein Sequence Database,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eur. J. Biochem. 23, 328-335, 1971
A;Title: Yeast thioredoxin. Amino-acid sequence around the
A;Reference number: 805793; MUID:72100583; PMID:4945270
A;Accession: S15360
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N,Alternate names: thioredoxine 2
C;Speciles: SchizosacCharomyces pombe
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
C;Accession: T39085; T46567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Keywords: redox-active disulfide F;2-103/Product: thioredoxin II #status predicted F;9-90/Domain: thioredoxin homology <THR>
                                                                                       A;Experimental source: strain 972h-; cosmid c7D4
R;Lenaers, G.; Perret, E.; Bonin, O.; Picard, A.; Caput,
submitted to the EMBL Data Library, October 1997
A;Description: TRX2, a fission yeast stress protein.
                                                                                                                                                                                                                                                                                                                                                                                                         R;Gentles, S.; Churcher, C.M.; Wood, V.; Barrell, submitted to the EMBL Data Library, September 199
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A;Cross-references: SGD:S0004033; MIPS:YLR043c
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                                                                                                                                                                                                                                                                                                          A;Status: preliminary; translated from GB/EMBL/DDB
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A; Status: preliminary; translated
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Pred. No. 1.3e+02;
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A; Dictus, Francis DNA
A; Molecule type: DNA
A; Residues: 1-103 <GLA>
A; Cross-references: UNIPROT: Q9S386; UNIPARC: UPI000005509E;
A; Cross-references: Strain Clip11262
                                                                                                                          D.; Jones, L.M.; Karst, U.S.; Dussurget, O.; E. Science 294, 849-852, 2001
Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.
A;Title: Comparative genomics of Listeria species,
A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Accession: AC1582
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                                                                                                                                                                                                                                                                          C;Species: Listeria innocua
C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 05-Oct-2004
C;Accession: AC1582
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A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.;
ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.;
A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
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C;Superfamily:
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A;Molecule type: DNA
A;Residues: 1-103 <GLA>
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                                                                                                               A; Status: preliminary
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Matches 4
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 Superfamily:
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thioredoxin; thioredoxin homology
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Pred. No. 1.3e+02;
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Pred. No. 1.3e+02;
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                                                                GB:AL592022; PIDN:CAC96427.1
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Voss, H.; Wehland
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Voss, H.; We
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Fsihi, H.
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Fsihi, H.
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R;Johnson, R.
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RESULT 27
B37192
thioredoxin - Bacillus subtilis
C;Species: Bacillus subtilis
C;Species: Bacillus subtilis
C;Date: 31-Jan-1992 #Bequence revision 26
C;Accession: B37192; H69726
R;Chen, N.Y.; Zhang, J.J.; Paulus, H.
J. Gen. Microbiol. 135, 2931-2940, 1989
                                                                                                                                                                                                                                                                                                                                                                                         R.Johnson, T.C.; Yee, B.C.; Carlson, D.E.; Buchanan, B.B.; Johnson, R.S.; Mathews, J. Bacteriol. 170, 2406-2408, 1988
A;Title: Thioredoxin from Rhodospirillum rubrum: primary structure and relation to A;Reference number: A28215; MUID:88198045; PMID:3129411
A;Accession: A28215
A;Accession: A28215
A;Molecule type: protein
A;Residues: 1-104 <-70Hs
A;Cross-references: UNIPROT:P10473; UNIPARC:UP10000171E24
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
F;7-90/Domain: thioredoxin homology <-THR>
F;29-32/Disulfide bonds: redox-active #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A,Title: Amino acid sequence of thioredoxin isolated from A,Reference number: A28086; MUID:88257078; PMID:3164311 A,Accession: A28086
A,Molecule type: protein A,Residues: 1-104 <JOH-A,Cross-references: UNIFROT:P08628; UNIPARC:UPI0000136E96 C,Superfamily: thioredoxin; thioredoxin homology C,Keywords: redox-active disulfide F;31-34/Disulfide bonds: redox-active #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Rhodospirillum rubrum
C;Date: 28-Aug-1989 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: A28215
C;Accession: A28215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            thioredoxin - Rhodospirillum rubrum C; Species: Rhodospirillum rubrum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; Species: Oryctolagus cuniculus (domestic rabbit)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9589-9597, 1988
sequence of thioredoxin isolated from
A28086; MUID:88257078; PMID:3164311
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Pred. No. 1.3
); Mismatches
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Pred. No. 1.3
0; Mismatches
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Pred. No. 1.3e+02;
; Mismatches 0;
                                                                        26-May-1994
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hes 0;
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A;Cross references UNIPROT:P14949; UNIPARC:UPI000006087B; GB:J03294; GB:M26384; NID:g14
A;Cross references UNIPROT:P14949; UNIPARC:UPI000006087B; GB:J03294; GB:M26384; NID:g14
C; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Chc
A; Ebrilich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Gallex
iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F.
Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois,
A;Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maucel
y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Fark, S.H.; Parro, V.; Pohl, T.M.; Portetelle
Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon,
A;Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Seror
akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama,
T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K.
A;Authors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.
A;Title: The complete genome sequence of the Gram, postiive bacterium Bacillus subtilis.
A;Accession: H69726
A.Scension: H69726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          thioredoxin I - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein G7746; protein YGR209c; thioredoxin
C;Species: Saccharomyces cerevisiae
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Eur. J. Biochem. 23, 328-335, 1971
A; Title: Yeast thioredoxin. Amino-acid sequence
A; Reference number: S05793; MUID:72100583; PMID:
                                                           A;Cross-references: UNIPARC:UPI000004F91C; R;Hall, D.E.; Baldesten, A.; Holmgren, A.; Eur. J. Biochem. 23, 328-335, 1971
                                                                                                                                                                                                                           J. Biol. Chem. 266, 9194-9202, 1991
A;Title: Thioredoxin deficiency in yeast prolongs S phase and A;Reference number: A39847; MUID:91225027; PMID:2026619
A;Accession: B39847
                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPROT:P22803; UNIPARC:UPI000004F91C; R;Muller, B.G.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              J. Biol. Chem. 266, 1692-1696, 1991
A;Title: Yeast thioredoxin genes.
A;Reference number: A38669; MUID:91107668; PMID:1988444
A;Accession: S15049
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A;Map position:
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A;Cross-references: UNIPARC:UPI000006087B; GB:Z99118; GB:AL009126; NID:g2635200; PIDN:CA
A;Experimental source: strain 168
C;Genetics:
                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-104 < MUL>
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A; Residues: 1-104 < GAN>
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; 9-90/Domain: thioredoxin homology <THR>
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Residues: 1-104 <CHE
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Accession: S15049; B39847; S05793; S53932; S61947; S64531; S63858; A38669
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Pred. No. 1.3e+02;
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                                                                                                GB:M62648; NID:g173049; Reichard, P.
                                    around the active-center disulfide
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A; Residues: 1-104 < GUW>
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A; Residues: 1-104 < GUE>
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A; Accession: S53932
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A;Cross-references: UNIPARC:UPI000004F91C; EMBL:Z49133; NID:g790489; A;Experimental source: strain S288C R;Song, J.M.; Cheung, E.; Rabinowitz, J.C. submitted to the EMBL Data Library, November 1995 A;Description: Analysis of the 15.6-kb fragment encompassing the ADE3 A;Reference number: S61947 A;Accession: S61947
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R;Guerreiro, P.; Barreiros, T.; Cyrne, L.; Soares, H.; Maia submitted to the Protein Sequence Database, May 1996
A;Reference number: S64517
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A;Residues: 1-104 <SON>
A;Cross-references: UNIPARC:UPI000004F91C; EMBL:U40843; NID:g1165213; PIDN:AAA85584.1;
C;Accession: S77780; S46921 R;Bork, P.; Ouzounis, C.; Casari, G.; Schneider, R.; Sander, C. Mol. Microbiol. 16, 955-967, 1995
A;Title: Exploring the Mycoplasma capricolum genome: a minimal A;Reference number: S77739; MUID:96059641; PMID:7476192
                                                                                                                              thioredoxin - Mycoplasma capricolum (fragment)
N;Alternate names: protein MC064
C;Species: Mycoplasma capricolum
C;Date: 09-Oct-1997 #sequence_revision 24-Oct-1997 #text_change 09-Jul-2004
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A;Residues: 1-104 <GUF>
A;Residues: 1-104 <GUF>
A;Cross-references: UNIPARC:UPI000004F91C; EMBL:Z49133; NID:g790489; PIDN:CAA89002.1;
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, April 1995
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A;Experimental source: strain S288C
R;Guerreiro, P.; Barreiros, T.; Soares, H.;
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A;Cross-references: SGD:S0003441; MIPS:YGR209c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL: Z72994; NID: g1323374; PIDN: CAA97236.1;
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                                                                                            R.; Sander, C.; Dolan, M.;
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A; Molecule type: DNA
A; Residues: 1-104 < KUR>
A; Cross-references: UNIPROT: Q97P68; UNIPARC: UPI00000519A5;
A; Experimental source: strain TIGR4
C; Genetics:
                                                                                                  thioredoxin-disulfide reductase (EC 1.8.1.9) [imported] - Streptococcus pneumoniae (;Species: Streptococcus pneumoniae C;Species: Streptococcus pneumoniae C;Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 09-Jul-2004 C;Accession: H98071
R;Hoskins, J.A.; Alborn Jr., W.; Arnold, J.; Blaszczak, L.; Burgett, S.; DeHoff, B. e, R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; McAhren, Y. P.; Sun, P.M.; Winkler, M.E.
J. Bacteriol. 183, 5709-5717, 2001
A;Authors: Yang, Y.; Young-Bellido, M.; Zhao, G.; Zook, C.; Baltz, R.H.; Jaskunas, A;Heference number: A97872; MUID:21429245; PMID:11544234
A;Accession: H98071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: redox-active disulfide F;13-93/Domain: thioredoxin homology <THR>
A;Residues: 1-104 <KUR>
A;Cross-references: UNI
C;Genetics:
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C;Superfamily:
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A;Accession: H95206
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Streptococcus pneumoniae
C;Date: 03-Aug-2001 #sequence_revision 03-Aug-2001 #text_change 09-Jul-2004
C;Accession: H95206
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A;Note: the nucleotide sequence was submitted to the EMBL Data Library, July 1994
C;Genetics:
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A;Molecule type: DNA
A;Residues: 1-104 <BOR>
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                                                                                    A;Status: preliminary
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                                                           A; Molecule type: DNA
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                    UNIPROT:Q8DNP9; UNIPARC:UPI00000E36AA;
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Pred. No. 1.3e+02;
); Mismatches 0;
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Pred. No. 1.3e+02;
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                    GB:AE007317; PIDN:AAL00405.1
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                                                                                                                                                                                                                thioredoxin 3 - slime mold (Dictyostelium discoideum)
(;Species: Dictyostelium discoideum)
C;Date: 28-Oct-1994 #sequence_revision 28-Oct-1994 #text_change 05-Oct-200
C;Accession: C46264
R;Wetterauer, B; Jacquot, J.P.; Veron, M.
J. Biol. Chem. 267, 9895-9904, 1992
A;Title: Thioredoxins from Dictyostelium discoideum are a developmentally A;Reference number: A46264; MUID:92250653; PMID:1577820
A;Accession: C46264
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C46264
                                                                                                  A;Status: preliminary; not compared with conceptual translation A;Molecule type: mRNA A;Molecule type: mRNA A;Residues: 1-104 <WET> A;Cross-references: UNIPROT:P29447; UNIPARC:UPI0000136D7B; GB:M91383; NID:g167932; C;Superfamily: thioredoxin; thioredoxin homology
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R;Reynolds, C.M.; Poole, L.B.; Hammel, K.E.; Wada, K.; Buchanan, submitted to the Protein Sequence Database, September 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Clostridium pasteurianum
C;Date: 02-Apr-2002 #sequence revision 02-Apr-2002 #text_change 05-Oct-2004
C;Accession: B59394; A59394
R;Reynolds, C.M.; Meyer, J.; Poole, L.B.
Biochemistry 41, 1990-2001, 2002
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A;Contents: annotation; purification and characterization C;Superfamily: thioredoxin; thioredoxin homology
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: protein
A; Residues: 1-104 < RE2>
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A; Accession: A59394
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C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: oxidoreductase
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    Clostridium pasteurianum

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Pred. No. 1.3e+02;
); Mismatches 0;
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Pred. No. 1.3e+02;
Score 31; DB 2;
Pred. No. 1.3e+02;
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R;Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, L.; ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Sekimizu, ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Hayashi, H.; Hiramatsu, K. C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K. Lancet 357, 1225-1240, 2001

A;Title: Whole genome sequencing of meticillin-resistant Stapylococcus aureus. A;Reference number: A89758; MUID:21311952; PMID:11418146

A;Accession: E89885
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A;Title: Complete genome sequence of the alkaliphilic bacterium A;Reference number: A83650; MUID:20512582; PMID:11058132
A;Accession: B84037
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-104 <570>
thioredoxin trxA [imported] - Bacillus halodurans (strain C-125)
C;Species: Bacillus halodurans
C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 05-Oct-2004
C;Accession: B84037
                                                                                                   RESULT
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A; Residues: 1-104 < KUR>
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C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 05-Oct-2004
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C;Superfamily: thioredoxin; thioredoxin homology
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                                                                                                                                                                                                                                                                                                                                                 ;Superfamily: thioredoxin; thioredoxin homology
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100.0%; Pred. No. 1.3e+02;
tive 0; Mismatches 0;
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Pred. No. 1.3e+02;
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                                     #text_change 05-Oct-2004
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  Query Match
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C;Accession: C81432
R;Parkhill, J.; Wren, B.W.; Mungall, K.; Ketley, J.M.; Churcher, C.; Basham, D.; Chilling; Quality, M.; Rajandream, M.A.; Rutherford, K.M.; VanVliet, A.; Whitehead, S.; Barrel Nature 403, 665-668, 2000
A;Title: The genome sequence of the food-borne pathogen Campylobacter jejuni reveals hyla;Reference number: A81250; MUID:20150912; PMID:10688204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 37
C81432
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A,Title: The complete genome sequence of the lactic acid bacterium Lactococcus lactis A,Reference number: A86625; MUID:21235186; PMID:11337471
A,Accession: D86630
A,Status: preliminary
                                                                             A;Residues: 1-105 <JON>
A;Residues: 1-105 <JON>
A;Cross-references: URIPROT:P08629; UNIPARC:UPI00001713BC;
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
E;9-92/Domain: thioredoxin homology <THR>
                                                                                                                                                                                    R;Jones, S.W.; Luk, K.C.
J. Biol. Chem. 263, 9607-9611, 1988
A;Title: Isolation of a chicken thioredoxin cDNA clone:
A;Reference number: A30006; MUID:88257080; PMID:2838473
A;Accession: A30006
A;Molecule type: mRNA
A;Molecule type: mRNA
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C;Species: Campylobacter jejuni
C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 05-Oct-2004
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A;Cross-references: UNIPROT:Q9CF37; UNIPARC:UPI0000C6AAD;
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C, Superfamily: thioredoxin; thioredoxin homology
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A; Residues: 1-104 < PAR>
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C;Superfamily: thioredoxin; thioredoxin homology
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                                                              F;32-35/Disulfide bonds:
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                                                                                                                                                                                                                                                                                                                                         C;Species: Gallus gallus (chicken)
C;Date: 31-Mar-1989 #sequence_revision
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A;Cross-references: UNIDARC:UPI000011065C; EMBL:X77584; NID:g453963; PIDN:CAA54687.1; PIR;WOllman, B.B.; d'Auriol, L.; Rimeky, L.; Shaw, A.; Jacquot, J.P.; Wingfield, P.; Grabe J. Biol. Chem. 263, 15506-15512, 1988
A;Title: Cloning and expression of a cDNA for human thioredoxin.
A;Reference number: A31993; MUID:89008454; PMID:3170595
A;Accession: A31993
                                                                                                                                                                                                                     A; Molecule type: mRNA
A; Residues: 1-105 < TAG2>
A; Cross-reference:
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C;Date: 30-Jun-1992 #secricon and a control of the control of 
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A;Title: Isolation and characterization of human thioredoxin-encoding genes
A;Reference number: JH0568; MUID:91340156; PMID:1874447
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R;Wetterauer, B.; Jacquot, J.P.; Veron, M.
J. Biol. Chem. 267, 9895-9904, 1992
A;Title: Thioredoxins from Dictyostelium discoideum are
A;Reference number: A46264; MUID:92250653; PMID:1577820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:P10599; UNIPARC:UPI000011065C; EMBL:X54539; NID:g37455; PIDN R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kondo, N.; Matsui, H.; Hamuro, J.; Brown, N.; Arai, EMBO J. 8, 757-764, 1989
A;Title: ATL-derived factor (ADF), an IL-2 receptor/Tac inducer homologous to thioredoxia, Reference number: S04106; MUID:89251607; PMID:2785919
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C;Superfamily: thioredoxin; thioredoxin homology
C;Reywords: redox-active disulfide
C;Reywords: redox-active disulfide
F;9-92/Domain: thioredoxin homology <THR>
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A;Contents: erratum
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Residues: 1-105 <TAG1>
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A;Reference number: A65534; PDB:1ERU
A;Contents: annotation; X-ray crystallography, 2.1 angstroms, oxidized for R;Forman-Ray, J.D.; Clore, G.M.; Gronenborn, A.M.
submitted to the Brookhaven Protein Data Bank, December 1990
A;Reference number: A50924; PDB:4TRX
A;Contents: annotation; conformation by (1)H-NMR, residues 1-73,'T',75-105
R;Forman-Kay, J.D.; Clore, G.M.; Wingfield, P.T.; Gronenborn, A.M.
Biochemistry 30, 2685-2698, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Note: described to be a surface-associated thioredoxin R;Wakasugi, H.; Rimsky, L.; Mahe, Y.; Kamel, A.M.; Fradelizi, D Proc. Natl. Acad. Sci. U.S.A. 84, 804-808, 1987
A;Title: Epstein-Barr virus-containing B-cell line produces an A;Reference number: A60870; MUID:87118252; PMID:3027706
A;Contents: annotation
R;Weichsel, A.; Gasdaska, J.R.; Powis, G.; Montfort, W.R. submitted to the Brookhaven Protein Data Bank, February 1996
A;Reference number: A65533; PDB:1ERT
                                                                                                                                                                                                                                                               A;Map position: 9q31-9q31
A;Introns: 8/3; 43/3; 63/3; 85/3
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A,Title: High-resolution three-dimensional structure of reduced recombinant human th A,Fatele: High-resolution three-dimensional structure of reduced recombinant human th A,Fateference number: A38953; MUID:91159399; PMID:2001356
A,Contents: annotation; conformation by (1)H- and (15)N-NMR
C,Comment: This small ubiquitous protein functions in many intracellular biological
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Biochem. J. 304, 861-867, 1994
A;Title: Characterization of a thioredoxin-related surface protein.
A;Reference number: S53453; MUID:95118305; PMID:7818492
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A;Molecule type: protein
A;Residues: 2-12, 'K',14-15,'XX',18-19,'X',21-22 <SIL>
A;Cross-references: UNIPARC:UPI0000171E1A
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A;Title: Human eosinophil cytotoxicity-enhancing factor.
A;Reference number: A60749; MUID:89309777; PMID:2745979
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Biochem. Biophys. Res. Commun. 175, 123-128, 1991
Biochem. Biophys. Res. Commun. 175, 123-128, 1991
A;Tille: Identification of a thioredoxin-related protein associated with plasma membran
A;Reference number: PT0079; MUID:91151337; PMID:1998498
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A;Residues: 1-21;38-57 <DEA>
A;Cross-references: UNIPARC:UPI00000353C5; UNIPARC:UPI0000171E1C
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R/Silberstein, D.S.; Ali, M.H.; Baker, S.L.; David,
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A;Cross-references: UNIPARC:UPI000003066B; GB:J04026; NID:g339648; PIDN:AAA74596.1;
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                                                                                                                     F;32-35/Disulfide
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                                                                                                                                                      ;Keywords: redox-active disulfide;2-105/Product: thioredoxin #status experimental <MAT>;9-92/Domain: thioredoxin homology <THR>
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genetics:
                                                                                                                     bonds:
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                                        Length 105;
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human thiore

path

residues

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N;Alternate names: ATL-derived factor (ADF)
C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Date: 21-Nov-1993 #sequence_revision 17-Oct-1997 #text_change 05-Oct-2004
C;Accession: JC4068; S44376; S04107
R;Matsui, M.; Taniguchi, Y.; Hirota, K.; Taketo, M.; Yodoi, J.
Gene 152, 165-171, 1995
A;Title: Structure of the mouse thioredoxin-encoding gene and its processed A;Reference number: JC4068; MUID:95137382; PMID:7835695
A;Accession: JC4068
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Biochem. Biophys. Res. Commun. 183, 170-175, 1992
A;Title: Thioredoxin gene expression is transcriptionally up-regulated by retinol A;Reference number: JS0667; MUID:92181438; PMID:1543487
A;Accession: JS0667
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JS0667
                                                                                                                                            A,Molecule type: mRNA
A;Residues: 1-93,'N',94-96,'ALT',100-104,'S' <TAG2>
A;Cross-references: UNIPARC:UPI0000171E1E; GB:X77585
C;Comment: This small ubiquitous protein functions in
                                                                                                                                                                                                                                                        A;Cross-references: UNIPARC:UPI0000163891; EMBL:X77585; NID:g453971; PIDN:CAA54688.1; R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kondo, N.; Matsui, H.; Hamuro, J.; Brown, N.; Ara EMBO J. 8, 757-764, 1989
A;Title: ATL-derived factor (ADF), an IL-2 receptor/Tac inducer homologous to thioredo A;Reference number: S04106; MUID:89251607; PMID:2785919
A;Accession: S04107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kando, N.; Matsui, EMBO J. 13, 2244, 1994
A;Reference number: S44375; MUID:94244626; PMID:8187776
A;Contents: erratum
A;Accession: S44376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:P29451; UNIPARC:UPI000016C493; GB:M84643; NID:g342338; PIDN C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: redox-active disulfide C;Keywords: redox-active disulfide F;9-92/Domain: thioredoxin homology <THR>
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C;Date: 30-Jun-1992 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: JS0667
A;Map position: 4:24.6
A;Introns: 29/2; 44/1; 84/2
C;Superfamily: thioredoxin; thioredoxin homology
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A; Residues: 1-105 < MAT>
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                                                                                                      A; Gene: MGI:Txn
                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-105 < TAG1>
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R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kando, N.; Matsui, H.; Hamuro, J.; Brown, N.; Arai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    thioredoxin -
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                                                                           A; Cross-references: MGI:36258
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Pred. No.
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hes 0;
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     F;30-33/Disulfide bonds: redox-active
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A;Molecule type: protein
A;Residues: 1-105 <MEN>
A;Residues: 1-105 <MEN>
A;Residues: 1-105 <MEN>
A;Cross-references: UNIPROT:P00275; UNIPARC:UPI0000136D69
A;Note: the source was designated as Corymebacterium nephridii
C;Comment: Thioredoxins are ubiquitous small hydrogen carrier proteins that participate
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                           C;ACCEBBAIN. AND C:
R;Meng, M.; Hogenkamp, H.P.C.
J. Biol. Chem. 256, 9174-9182, 1981
A;Title: Purification, characterization, an
A:Peference number: A00281; MUID:81264365;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: UNIPROT:P11232; UNIPARC:UPI000003066A; EMBL:X14 R;Dean, M.F.; Martin, H.; Sansom, P.A. Biochem. J. 304, 861-867, 1994
A;Title: Characterization of a thioredoxin-related surface protein. A;Reference number: S53453; MUID:95118305; PMID:7818492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Rattus norvegicus (Norway rat)
C;Date: 31-Mar-1990 #sequence revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: S04352; S66372
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S04352
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A;Note: 12-Lys, 15-Gly, 16-Leu, 17-Gln, and 18-Leu were also
A;Note: described to be a surface-associated thioredoxin
C;Superfamily: thioredoxin; thioredoxin homology
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C; Keywords: redox-active disulfide F;8-91/Domain: thioredoxin homology <THR>
                                                                                                                                                                                                                                                                                                                                                                                    thioredoxin - coryneform bacterium ATCC11425
C;Species: coryneform bacterium ATCC11425
C;Date: 02-Apr-1982 #sequence_revision 02-Apr-1982 #text_change 05-Oct-2004
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                                                                                                                                                                                                                          A; Accession: A00281
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A; Residues: 1-105 < TON>
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5; PMID:7021558
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PID

#status experimental

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C;Species: Synechocystis sp.
A;Variety: PCC 6803
C;Date: 25-Apr-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
C;Date: 25-Apr-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
C;Accession: S76386
R;Kaneko, T.; Sato, S.; Kotani, H.; Tanaka, A.; Asamizu, E.; Nakamura, Y.; Ro, Ckumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Watanabe, A.; Yamadi O, K.; Okumura, S.; Shimpo, Shimpo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Ogata, H.; Audic, S.; Renesto-Audiffren, P.; Fournier, P.E.; Barbe, V.; Samson, Science 293, 2093-2098, 2001
A;Title: Mechanisms of Evolution in Rickettsia conorii and Rickettsia prowazekii.
A;Reference number: A97700; MUID:21442074; PMID:11557893
A;Accession: B97700
A;Status: preliminary
RESULT
AG2042
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C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 05-Oct-2004
C;Accession: B97700
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C;Superfamily:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          thioredoxin M-1 - Synechocystis sp. N;Alternate names: protein slr0233
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Genetics:
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100.0%; Pred. No. 1.3e+02;
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Yamada, M.; Yasud
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RiPille, S.; Chuat, J.C.; Breton, A.M.; Cle
J. Bacteriol. 172, 1556-1561, 1990
A;Title: Cloning, nucleotide sequence, and
A;Reference number: A35135; MUID:90170874;
A;Accession: A35135
                                                                                                                                                                          C;Species: Rhodobacter sphaeroides
C;Date: 27-Jul-1990 #sequence_revision
C;Accession: A35135; A30508
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A35135
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J. Bacteriol. 183, 4823-4838, 2001
A;Title: Genome Sequence and Comparative Analysis of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rnioredoxin [imported] - Clostridium acetobutylicum
C;Species: Clostridium acetobutylicum
C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 05-Oct-2004
C;Accession: D97279
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A;Note: Nostoc sp. strain PCC 7120 is a synonym of C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 C;Accession: AG2042
A;Molecule type: DNA
A;Residues: 1-106 <PIL>
A;Cross-references: UNIPROT:P08058; UNIPARC:UPI000016FF78; GB:M33806; NID:g152044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.S.; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.
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C; Superfamily:
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A;Cross-referenceB: UNIPROT:Q8YVT1; UNIPARC:UPI00000CE247; GB:BA000019; PIDN:BAB73592.1.
A;Experimental source: strain PCC 7120
C;Genetics:
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A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium AnA;Reference number: AB1807; MUID:21595285; PMID:11759840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Gene: CAC3083
C;Superfamily: thioredoxin; thioredoxin homology
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A;Experimental source: Clostridium acetobutylicum ATCC824
                                                                                                                                                                                                                                            thioredoxin
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A;Residues: 1-105 <KUR>
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Nakazaki, N.; Shimpo, S.;

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    Rhodobacter sphaeroides

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Pred. No. 1.3e+02
); Mismatches 0
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Pred. No. 1.3e+02;
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PMID:2137818
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R;Clement-Metral, J.D.; Holmgren, A.; Cambillau, C.; Joernvall, F
Bux. J. Biochem. 172, 413-419, 1988
A;Title: Amino acid sequence determination and three-dimensional
A;Reference number: A30508; MUID:88166714; PMID:3280308
A;Accession: A30508
A;Molecule type: protein
A;Residues: 2-63, Z,65-106 <CLE>
A;Cross-references: UNIPARC;UPT00000171E23
                                                                                                           C;Accession: H64622; C71890

R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; Mcison, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujii, C.; Bowman, C.; Watthey Nature 388, 539-547, 1997

A;Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser A;Title: The complete genome sequence of the gastric pathogen Helicobacter pylori. A;Reference number: A64520; MUID:97394467; PMID:9252185

A;Accession: H64622
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C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
F;9-92/Domain: thioredoxin homology <THR>
F;31-34/Disulfide bonds: redox-active #status pr
 A;Experimental source: strain 26695 R;Alm, R.A.; Ling, L.S.L.; Moir, D.
                                                                                                                                                                                                                                                                                                         C; Species: Helicobacter pylori
A; Variety: strains J99, 26695
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A; Residues: 1-106 < LAB>
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A; Accession: $33357
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R;Labudova, O.; Nemethova,
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                                                       A; Molecule type: DNA
A; Residues: 1-106 < TOM>
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                                     A; Cross-references: UNIPROT: P56430; UNIPARC: UPI0000136E82;
                                                                                             A;Status: nucleic acid sequence
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Best Local S
Matches 4
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;9-92/Domain: thioredoxin homology <THR>
;31-34/Disulfide bonds: redox-active #status predicted
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Species: Streptomyces aureofaciens
Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change
Accession: S33357
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100.0%; Pred. No.
Moir,
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Library, April 1993
of a protein-coding part of the
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Pred. No. 1.3e+02;
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 P.C.; Smith, D.R.
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A;Cross-references: UNIPROT:Q8UJA6; UNIPARC:UPI0000164451; GB:AE008688; PIDN:AAL41053.1
A;Experimental source: strain C58 (Dupont)
C;Genetics:
                                                                                                                 A; Reference number: A; Accession: AG2579
                                                                                                                                                       ster, E.W.
A; Title: The Genome of
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A; Residues: 1-106 < COH>
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                                                                                             A;Status: preliminary
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C;Accession: AG2579

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R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo, erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClel; Karp, P.; Romero, P.; Zhang, S.
Science 294, 2317-2323, 2001

A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Genomic sequence comparison of two A;Reference number: A71800; MUID:99120557; I A;Accession: C71890
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Cohen, G.; Argaman, A.; Schreiber, R.; Mislovati, M.; Aharonowitz, J. Bacteriol. 176, 973-984, 1994
A;Title: The thioredoxin system of Penicillium chrysogenum and its p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Superfamily: thioredoxin; thioredoxin homology F;10-92/Domain: thioredoxin homology <THR>
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C;Accession: A49888; S38886
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Ives, C.; Gibson, R.; Merberg, D.; Mills, S.D.; Jiang, Q.; Nature 397, 176-180, 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Cross-references: UNIPROT: P34723; UNIPARC: UPI0000136E91;
                                                                                                                                                                                                 thioredoxin; thioredoxin homology
of the Natural Genetic Engineer Agrobacterium tumefaciens AB2577; MUID:21608550; PMID:11743193
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Pred. No. 1.3e+02;
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A;Cross-references: UNIPARC:UPI00001655B5; GB:M14736; NID:g142117; PIDN:AAA22049.1;
R;Gleason, F.K.; Whittaker, M.M.; Holmgren, A.; Joernvall, H.

J. Biol. Chem. 260, 9567-9573, 1985
A;Title: The primary structure of thioredoxin from the filamentous cyanobacterium An A;Reference number: A23910; MUID:85261357; PMID:3926769
A;Accession: A23910
A;Molecule type: protein
A;Residues: 2-107 <GLE>
A;Residues: 2-107 <GLE>
A;Residues: 2-107 <GLE>
A;Resperimental source: PCC 7119, ATCC 29151
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: heat-stable protein; redox-active disulfide
F;2-107/Product: thioredoxin #status experimental <MAT>
F;10-93/Domain: thioredoxin homology <THR>
F;32-35/Disulfide bonds: redox-active #status predicted
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A;Residues: 1-107 <MUL>
A;Residues: 1-107 <MUL>
A;Cross-references: UNIPARC:UPI0000001823; GB:J04475; NID:g142153; PIDN:AAA22057.1; PID:A;Note: the source is designated as Anacystis nidulans R2, which is also called Synechoo C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Muller, B.G.D.; Buchanan, B.B.
J. Biol. Chem. 264, 4008-4014, 1989
A;Title: Thioredoxin is essential for photosynthetic growth.
A;Reference number: A32956; MUID:89139466; PMID:2492995
A;Accession: A32956
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A32956
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C;Speciles: Synechococcus sp.
C;Date: 17-Jul-1992 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: A32956; A30842
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A;Map position: circular chromosome
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-107 < RES>
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Best Local S
Matches 4
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Dec-1988 #sequence_revision 11-Apr-1997 #text_change 05-Oct-2004
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Pred. No.
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A;Gene: trx
C;Superfamily: thioredoxin; thioredoxin homology
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
F;10-93/Domain: thioredoxin homology <THR>
F;32-35/Disulfide bonds: redox-active #status pr
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A;Title: Physical map of the plastid genome of the unicellular A;Reference number: $45488; MUID:95094309; PMID:8001167
A;Accession: $45488
                                                                                                                                                                                                             A;Molecule type: DNA
A;Residues: 'M',6,'IDEL',11-12,'DT',15-16,'LQ',19,'HQ',22-38,'A',40,'IL',43-44,'I',46-4
A;Cross-references: UNIPARC:UPI00000A275B
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A;Residues: 1-107 <JOH>
A;Cross-references: UNIPROT:P09857; UNIPARC:UPI0000136E76
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Biochemistry 26, 1209-1214, 1987
Biochemistry 26, 1209-1214, 1987
A;Tille: The primary structure of thioredoxin from Chromatium vinosum
A;Reference number: A26622; MUID:87185419; PMID:3567166
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C;Species: Cyanidium caldarium
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C;Date: 31-Mar-1988 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
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21
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Conservative

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Indels

0

Gaps

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C;Species: Drosophila melanogaster
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 05-Oct-2004
C;Accession: S47867
R;Salz, H.K.; Plickinger, T.W.; Mittendorf, E.; Pellicena-Palle, A.; Petschek, J.P.; Brogenetics 136, 1075-1086, 1994
A;Title: The Drosophila maternal effect locus deadhead encodes a thioredoxin homolog req
A;Reference number: S47867; MUID:94274010; PMID:7516301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               thioredoxin A - red alga (Porphyra purpurea) chloroplast
C;Species: chloroplast Porphyra purpurea
C;Date: 19-Mar-1997 #sequence_revision 09-May-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          F;31-34/Disulfide bonds:
                      RESULT 60
S46521
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S73146
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A; Residues: 1-107 < SAL>
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                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-107 < REI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plant Mol. Biol. Rep. 13, 333-335, 1995
A;Title: Complete nucleotide sequence of the Porphyra purpurea chloroplast genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R; Reith, M.; Munholland, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Gene: FlyBase:dhd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:P47938; UNIPARC:UPI0000124A52; EMBL:L27072; NID:g435591; PII
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  thioredoxin -
                                                                                                                                                                                                                                                                                                                            A; Genome: chloroplast
                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Status: preliminary; nucleic acid sequence not shown; translation not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Accession: S73146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Plant Mol.
                                                                                                                                                                        Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                  Superfamily: thioredoxin; thioredoxin homology; Keywords: chloroplast; redox-active disulfide; 9-92/Domain: thioredoxin homology <THR>
                                                                                                                                                                                                                                                                                                                                                   ;Gene: trxA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Accession: S73146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Reywords: redox-active disulfide;8-92/Domain: thioredoxin homology <THR>
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                                                                                                CGPC 34
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Porphyra yezoensis chloroplast
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Conservative
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Pred. No.
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Pred. No.
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  A;Molecule type: DNA
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C;Species: chloroplast Porphyra yezoensis
C;Date: 19-Mar-1997 #sequence_revision 06-Jun-1997 #text_change 09-Jul-2004
C;Accession: $46521
R;Reynolds, A.E.; Chesnick, J.M.; Woolford, J.; Cattolico, R.A.
Plant Mol. Biol. 25, 13-21, 1994
A;Title: Chloroplast encoded thioredoxin genes in the red algae Porphyra yezoensis and G
A;Reference number: $46521; MUID:94272009; PMID:8003693
A;Accession: $46521
A;Status: preliminary
A;Kolecule type: DNA
A;Residues: 1-107 <REY>
A;Cross-references: UNIPROT:P50254; UNIPARC:UPI0000136E94; EMBL:X76612; NID:g509276; PII
C;Genetics:
A;Genome: chloroplast
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: chloroplast; redox-active disulfide
F;9-92/Domain: thioredoxin homology
C;Keynords: thioredoxin homology
                                                                                              C;Species: Synechocystis sp.
A;Variety: PCC 6803
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995
C;Accession: S46958; S'6679
R;Navarro, F.; Florencio, F.J.
                                                                                                                                                                                                        thioredoxin A - Synechocystis sp. N/Alternate names: protein slr0623 C/Species: Synechocystis sp.
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A;Introns: 31/2
C;Superfamily: thioredoxin; thioredoxin homology
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A;Residues: 1-107 <WOB>
A;Crosa-references: UNIPROT:Q9TXY8; UNIPARC:UPI0000077021; EMBL:AF106589; PIDN:AAC78230
A;Experimental source: strain Bristol N2; clone Y44E3A
C;Genetics:
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submitted to the EMBL Data Library, November 1998
A;Description: The sequence of C. elegans cosmid Y44E3A.
A;Reference number: Z21422
submitted to the EMBL Data Library, July 1994
A;Description: The gene encoding for thioredo:
A;Reference number: S46957
A;Accession: S46958
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C;Accession:
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F;31-34/Disulfide bonds: redox-active #status predicted
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Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change
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Best Local
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Best Local Similarity
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100.0%; Pred. No. 1.3e+02;
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100.0%; Pred. No. 1.3e+02;
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                                                      thioredoxin
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                                                      the
                                                      cyanobacterium Synechocy
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A; Gene: TRXRP1
A; Map position:
C; Superfamily:
C;Species: Streptomyces clavuligerus
C;Date: 08-Sep-1995 #sequence_revision 08-Sep-1995 #text_change 05-Oct-2004
C;Accession: B53307
R;Cohen, G: Yanko, M: Mislovati, M: Argaman, A: Schreiber, R: Av-Gay, 'Y:; I
Bacteriol. 175, 5159-5167, 1993
A;Title: Thioredoxin-thioredoxin reductase system of Streptomyces clavuligerus:
A;Reference number: A53307; MUID:93352422; PMID:8349555
A;Accession: B53307
A;Status: preliminary
A;Molecule type: DNA
                                                                                                                                                                                                                                                                   RESULT 64
B53307
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A;Residues: 1-107 <NAV>
A;Cross-references: UNIFROT:P52231; UNIPARC:UPI0000164C7C; EMBL:x80486; NID:g515939; PII R;Kaneko, T.; Sato, S.; Kotani, H.; Tanaka, A.; Asamizu, E.; Nakamura, Y.; Miyajima, N.; C,; Kaneko, T.; Sato, S.; Shimpo, S.; Takeuchi, C.; Wada, T.; Watanabe, A.; Yamada, M.; Yasud DNA Res. 3, 109-136, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Leishmania major

C;Date: 24-Mar-1999 #sequence_revision 24-Mar-1999 #text_change 09-Jul-2004

C;Accession: B81458; T02814

C;Accession: B91458; T02814

R;Myler, P.J.; Audleman, L.; deVos, T.; Hixson, G.; Kiser, P.; Lemley, C.; Magness, C.;

Proc. Natl. Acad. Sci. U.S.A. 96, 2902-2906, 1999

A;Title: Leishmania major
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T02814
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C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
E;10-93/Domain: thioredoxin homology <THR>
                                                                                                                                                                                     thioredoxin - Streptomyces clavuligerus C;Species: Streptomyces clavuligerus C;Date: 08-Sep-1995 #sequence_revision
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-107 < K
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Best Local S
Matches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      position:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 31; DB 2; 100.0%; Pred. No. 1.3e+02; o. Mismatches 0;
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CACCESSION: AH3504
CACCESSION: AH3504
R;DelVecchio, V.G.; Kapatral, V.; Redkar, R.J.; Patra, G.; Mujer, C.; Los, T.;
R;DelVecchio, V.G.; Kapatral, V.; Redkar, R.J.; Patra, G.; Mujer, C.; Los, T.;
Mazur, M.; Goltsman, E.; Selkov, E.; Elzer, P.H.; Hagius, S.; O'Callaghan,
Proc. Natl. Acad. Sci. U.S.A. 99, 443-448, 2002
A;Title: The genome sequence of the facultative intracellular pathogen Brucel.
A;Reference number: AD3252; PMID:11756688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Residues: 1-107 <COH>
A;Cross-references: UNI
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. strain PCC 7120
C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change 05-Oct-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Start codon: GTG
C;Superfamily: thioredoxin;
F;11-93/Domain: thioredoxin
                                                                                                                                                       A;Cross-references: UNIPROT:Q8YE56; UNIPROT:Q8FXY9; UNIPARC:UPI00000582C9; GB:AE008917; A;Experimental source: strain 16M
                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-107 < KUR>
                                                                                                                                                                                                                                                                                                                                                                                       C;Species: Brucella melitensis
C;Date: 01-Feb-2002 #sequence_revision 01-Feb-2002 #text_change 05-Oct-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPROT:P06544; UNIPARC:UPI00001655B5; GB:BA000019; PIDN:BAB77576.1 A;Experimental source: strain PCC 7120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nakazaki, N.; Shimpo, S.;
DNA Res. 8, 205-213, 2001
A;Title: Complete Genomic
                                                                                                                                                                                                                                 A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                     thioredoxin C-1 [imported] - Brucella melitensis (strain 16M)
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A; Residues: 1-107 < KUR>
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                                                                           ;Superfamily: thioredoxin; thioredoxin homology
                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , Y.; Wolk, C.P.; Kuritz, T.;
S.; Sugimoto, M.; Takazawa, N
                  100.0%;
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7; MUID:21595285; PMID:11759840
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Score 31; DH
Pred. No. 1.3
D; Mismatches
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Pred. No.
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M.; Yamada,
                                  Length 107
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Similarity 4; Conserv

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RiFleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.P.; Kerlavage, Gocayne, J.D.; Scott, J.; Shirley, R.; Liu, L.I.; Glodek, A.; Kelley, J.M.; Weidman, D.M.; Brandon, R.C.; Fine, L.D.; Fritchman, J.L.; Fuhrmann, J.L.; Geoghagen, N.S.M. Science 269, 496-512, 1995
A;Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Venter, A;Title: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd.
A;Reference number: A64000; MUID:9535039; PMID:7542800
A;Accession: E64047
A;Status: nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nature 406, 959-964, 2000
A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic pathc A;Reference number: A82950; MUID:20437337; PMID:10984043
A;Accession: G82991
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A;Note: named as homolog to a protein from Anacystis nidulans
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
F;10-93/Domain: thioredoxin homology <THR>
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R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig,
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C;Date: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 05-Oct-2004
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  thioredoxin -
                           RESULT 69
B55124
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                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-108 <STO>
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C;Species: Pseudomonas aeruginosa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Status: preliminary
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Best Local
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thioredoxin 1 [imported] - Yersinia pestis (strain CO92)
C;Species: Yersinia pestis
C;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 05-Oct-2004
C;Accession: AD0471
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A;Note: unidentified residues are Ile or Leu
A;Note: the species is identified as Chlorobium thiosulfatophilum
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: photosynthesis; redox-active disulfide
F;10-93/Domain: thioredoxin homology <THR>
F;32-35/Disulfide bonds: redox-active #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Parkhill, J.; Wren, B.W.; Thomson, N.R.; Titball, R.W.; Holden, M.T.G.; Prentice, deno-Tarraga, A.M.; Chillingworth, T.; Cronin, A.; Davies, R.M.; Davis, P.; Dougan, il, M.; Rutherford, K.; Simmonds, M.; Skelton, J.; Stevens, K.; Whitehead, S.; Barre Nature 413, 523-527, 2001
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A;Residues: 1-15,'XX',18-41,'X',43-44,'X',46-53,'XX',56-65,'X',67-74,'X',76-78,'XX',81-9
A;Cross-references: UNIPARC:UPI0000174DED
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A;Accession: A27801
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A;Residues: 1-108 <BIE>
A;Cross-references: UNIPARC:UPI0000174DEC
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C;Superfamily: thioredoxin; thioredoxin homology
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C;Genetics:
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A; Residues: 1-108 < KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Title: Genome sequence of Yersinia pestis, the causative A;Reference number: AB0001; MUID:21470413; PMID:11586360
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Matches
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InterPro; IPR007958; toxin 5.

Diam; PF05294; Toxin 5; 1.

Direct protein sequencing; Neurotoxin; Toxin.

DISULFID 18 By similarity.

DISULFID 4 25 By similarity.

DISULFID 19 30 By similarity.

DISULFID 19 32 By similarity.

SEQUENCE 35 AA; 3673 MW; 213E69262289EB5A CRC64;
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21-JUL-1986 (Rel.
05-JUL-2004 (Rel.
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Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida;
Buthida; Buthoidea; Buthidae; Androctonus.
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"Characterization of
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    -!- SUBCELLULAR LOCATION: Secreted.
    -!- TISSUE SPECIFICITY: Expressed by the venom gland.
    -!- SIMILARITY: Belongs to the short scorpion toxin family.

                                                                                                                                                                                                                                                                                                                                                                                                           Toxicon 23:113-125(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=85193276; PubMed=3992595; DOI=10.1016/0041-0101(85)90114-X;
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Q6UWD8_HUMAN
Q7K4P2_DROME
Q8C708_MOUSE
Q8C708_MOUSE
Q5BK39_RAT
Q6P3A6_MOUSE
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Q6SN76_SPR.IM
Q6SNH6_COLGU
Q6SNP3_PAPPA
Q8LDI1_ARATH
Q6A5V5_PROAC
Q8RG01_FUSINN
Q9CB61_MYCLE
Q6CER1_MYCLE
Q6CER4_ORYSA
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six of which are toxic
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Q8gmh2
Q9gnf4
Q9z234
Q8nab0
Q6nwd8
Q7k4p2
Q8r670
Q5bk39
Q6p3a6
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Q6sn76
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Q6snp3
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Q6a5v5
Q8cg01
                                                                                                                                                                                                                                                                                                   restrictions
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arabidopsis
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BS14 BUTSI
P59887;
10-OCT-2003
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01-JUN-1998
01-MAR-2003
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DISULFID
DISULFID
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Chloride
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Voelter W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      use
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Buthida; Buthoidea;
                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
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Tapirus.
NCBI_TaxID=9802;
                                                                                                                                                                                                                                                                                                                                                                                                                                           DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro;
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RNXCOCC
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10-OCT-2003 (Rel. 42, Last sequence up:
05-JUL-2004 (Rel. 44, Last annotation:
Chlorotoxin-like peptide Bs 14 (Bs14).
Buthus sindicus (Scorpion).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN the European Bioinformatics Institute. There are no rest
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  from scorpion (Buthus sindicus) venom.";
Comp. Biochem. Physiol. 121A:323-332(1998).
-!- FUNCTION: Probable chloride channel lig
                                            Luteinizing hormone beta-subunit (Fragment).

Name=LH-beta;
Tapirus indicus (Asiatic tapir) (Malayan tapir).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;

Mammalia; Eutheria; Laurasiatheria; Perissodactyla;
                                                                                                                                                                                            046621 TAPIN PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Purification and primary structure of low
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NOTE=Ref.1.
SIMILARITY: Belongs to the s
channel inhibitor subfamily.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      conductance chloride channels (By similarity).
SUBCELLULAR LOCATION: Secreted.
TISSUE SPECIFICITY: Expressed by the venom gland
MASS SPECTROMETRY: MW=3848.3; METHOD=MALDI; RANG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PF05294; Toxin 5; 1.

PF05294; Toxin 5; 1.

ide channel inhibitor; Neurotoxin; Toxin

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Conservative
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                                                                                                              (TYEMBLrel. 06, Created)
(TYEMBLrel. 06, Last sequence update)
(TYEMBLEL. 23, Last annotation update)
hormone beta-subunit (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                   1 inhibitor; 1
1 18
4 25
15 30
19 32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PubMed=10048185; DOI=10.1016/S1095-6433(98)10140-X;
S., Schuetz J., Kayed R., Abbasi A., Zaidi Z.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AND MASS
                                                                                                                                                                                                                                                                                                                                                                                                                      3847
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Buthidae; Mesobuthus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               toxin_5.
                                                                                                                                                                                                                                                                                                                                                                    . 0 %;
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                                                                                                                                                                                                                                                                                                                                                                                                                  By similarity.
By similarity.
By similarity.
By similarity.
S36117F8E50165A0
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                                                                                                                                                                                                                                                                                                                                                    Score 31; DE Pred. No. 2.1); Mismatches
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                                                Euteleostomi;
Tapiridae;
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01-JUN-1998
01-JUN-2003
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NON_TER
SEQUENCE
                                                                                   01-NOV-1996 (TrEMBLrel. 01, 01-NOV-1996 (TrEMBLrel. 01, 01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
Fischer S., Veits J., Meyer
Submitted (FEB-1998) to the
EMBL; AP047605; AAC04363.1;
HSSP; P01233; 1HCN.
                             Name=Mst87Fa; Synonyms=DhTc3;
Drosophila hydei (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Inse
Neoptera; Endopterygota; Diptera; Brachycera;
                                                                                                                  JT 5

17 DROHY

023947 DROHY PRELIMINARY;

023947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GO; GO:0005179; F:hormone activity; IEA.
InterPro; IPR006208; Cys_knot.
Pfam; PF00007; Cys_knot; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fischer S., Veits J., Meyer H.H.D.;
Submitted (FEB-1998) to the EMBL/GenBank/DDBJ
EMBL; AF047606; AAC04364.1; -; Genomic_DNA.
HSSP; P01233; 1HCN.
         Ephydroidea; Drosophilidae; Drosophila
NCBI_TaxID=7224;
                                                                        Testis-specific RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-JUN-1998 (TrEMBLrel. 06, Created)
01-JUN-1998 (TrEMBLrel. 06, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last sanotation update)
Luteinizing hormone/chorionic gonadotrophin beta-subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        046620
                                                                                                                                                                                                                                                                                                          Chorion.
                                                                                                                                                                                                                                                                                                                   Pfam; PF00007;
                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=73336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Equus asinus somalicus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=LH/CG-beta;
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0007; Cys_knot; 1.
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                                                                                                                                                                                                                                                                         4332 MW;
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Laurasiatheria; Perissodactyla;
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Last sequence
Last annual
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EMBL/GenBank/DDBJ
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Pred.
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Pred. No. 2.2e+0
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                                                                                     sequence update) annotation updat
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                                         Insecta;
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                                Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                    databases
                                                                                                                                                                                                                                                 Length 42;
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                                         Pterygota;
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Equidae; Equu
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RESULT 7

QBMLNS_DROME
QBMLNS_DROME PRELIMINARY;
AC QBMLNS;
DT 01-0CT-2002 (TrEMBLrel. 22,
DT 01-0CT-2002 (TrEMBLrel. 31,
DE CG30430-PA (IP04475p).
GN Name=CG30430; ORFNames=CG304
OS Drosophila melanogaster (Fro
CC Eukaryota; Metazoa; Arthropy
CC Neoptera; Endopterygota; Did
Ephydroidea; Drosophilidae;
OX NCBI_TaxID=7227;
RN [1]
RA Adams M.D., Celniker S.E., Is
RA George R.A., Lewis S.E., R;
RA Brandon R.C., Rogers Y.-H.C
RA Wan K.H., Doyle C., Baxter J
RA Adams M.D., E., Agbayani A., An
RA Ballew R.M., Basu A., Baxen
RA Beeson K.Y., Benos P.V., Be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q23948 DACHY
ID Q23948;
AC Q23948;
DT 01-NOV-1
DT 01-MAR-2
DT 10-MAR-2
DT 10-MAR-2
DE Testis-e
GN Name=Mst
OS Drosophi
OC Eukaryot
OC Neopters
OC Ephydroi
OC Ephydroi
OC Ephydroi
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RP NUCLEOTI
RA Kuech A.
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NUCLEOTIDE SEQUENCE.

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Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,

Adams M.D., Celniker S.E., Holt R.A., Hoskins R.A., Galle R.F.,

George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,

Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,

Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,

Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,

Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,

Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,

Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
                                                                                                                                                                                                                                                                           Name=CG30430; ORFNames=CG30430;
Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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Submitted (DEC-1990) to the EMBL/GenBank/DDBJ
EMBL; X58114; CAA41118.1; -; Genomic_DNA.
FlyBase; FBgn0015097; Dhyd\Mst87Fb.
SEQUENCE 49 AA; 4550 MW; 24758FEC51369F1D
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Submitted (DEC-1990) to the EMBL/GenBank/DDBJ databases.
EMBL; X58114; CAA41117.1; -; Genomic_DNA.
FlyBase; FBgn001096; Dhyd\Mst87Fa.
SEQUENCE 49 AA; 4610 MW; BE158FEC51369F01 CRC64;
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01-OCT-2002 (TremBirel. 22,
13-SEP-2005 (TremBirel. 31,
CG30430-PA (IP04475p).
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Name=Mst87Fb; Synonyms=DhTc2;
Drosophila hydei (Fruit fly).
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01-NOV-1996 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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Q23948;
Q1-NOV-1996 (TrEMBLrel. 0:
01-NOV-1996 (TrEMBLrel. 0:
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NCBI_TaxID=7224;
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Pred. No. 2.7e+02;
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Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Center A., Chandra I.,
RA de Pablos B., Delcher A., Deng Z., Mays A.D., Davies P.,
RA Dodoon K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kemison J.A., Ketchum K.A.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulip D., Lai Z.,
RA Liako P., Lei Y., Levitsky A.A., Li J.H., Li Z., Liang Y., Lin X.,
RA Mount S.M., Moy M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,
RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Williams S.M., Woodage T., Worley E., Wang S., Yao Q.A.,
RA Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zhang Y.H., Myers E.M., Rushin G.M., Pann G., Zhan G., Zhao Q., Zhang L.,
RA Spier E., Spradling A.N., Zhong W., Zhang G., Zhao Q., Zhang C., Schence 287:2185-2195(2000).
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Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
                                                                                                                                                                                                                                                                                                                                                        MEDLINE=22426069; PubMed=12537572;
MiBra S., Crosby M.A., Mungall C.J., Malthews B.B., Campbell K.S.,
Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E.
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.O.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                    Berkeley Drosophila Genome Project;
Celniker S., Carlson J., Wan K., Pf
Hoskins R., Stapleton M., Pacleb J.
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Kaminker J.S., Bergman C.M., Kronmiller
Patel S., Frise E., Wheeler D.A., Lewis
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EMBL/GenBank/DDBJ databases
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01-JUL-1993 (Rel
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PROSITE; P
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InterPro; IPR001450; 4Fe4S ferredoxin.
InterPro; IPR006081; Defensin_alpha.
InterPro; IPR001007; VMF C.
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"Association of misexpression with sterility in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ephydroidea; Dro:
NCBI_TaxID=7226;
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EMBL; BT023672; AAY85072.1; -; mRNA.
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Ephydroidea; Drosophilidae;
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My549956; AAS55565.1; -; Genomic DNA.
GO:0005576; C:extracellular region; IEA.
GO:0042742; P:defense response to bacteria;
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PS00269; DEFENSIN; 1.
PS01208; VWFC_1; 1.
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Pred. No. 2.8
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update)

Name=Mst84Dc; ORFNames=CG17945;

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RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Golle R.P., RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Golle R.P., RA Adams M.D., Celniker S.E., Richards S., Ashburner M., Pfelffer B.D., RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., RA Burton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., RA Burton G.G., Rogers Y.-H.C., Blazef R.G., Champe M., Pfelffer B.D., Ra Maril J.F., Agbayani A., An H.-J., Andrews Pfelmhoch C.R., Miklos G.I.G., RA Abril J.F., Agbayani A., An H.-J., Andrews Pfelmhoch C., Baldwin D., Ballow R.M., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Ra Beson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Ra Burtis K.C., Busam D.A., Butler H., Cadleu E., Center A., Chandra I., Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., Ra Dorkova D., Bolcher A., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M., Posler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.R., Cholos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M., Posler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.R., Alaris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J., Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ilegwam C., Lasko P., Lei Y., Leviteky A.A., Li J.H., Li Z., Liang Y., Lin X., Liang Y., Lin X., Liang Y., Lin X., Liang Y., Lin X., Liang Y., Matcei B., McIncosh T.C., McLeod M.P., McPherson D. D., Li Y., Leviteky A.A., Li J.H., Li Z., Liang Y., Lin X., Liang Y., Matcei B., McIncosh T.C., McLeod M.P., McPherson D.L., Nelson D.R., Nelson D.L., Murphy B., Murphy L., Muzny D.M., Nelson D.L., Ra Melson D.R., Nelson S., Sanders R.D.C., Scheeler F., Shen H., She P., Shen H., She P., She P., She H., Wang X., Smith H.O., Mang Y., Wang X., She P., 
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MEDLINE=92102953; PubMed=1684716; DOI=10.1016/0925-4773(91)90064-D;
MEDLINE=92102953; PubMed=1684716; DOI=10.1016/0925-4773(91)90064-D;
Kuhn R., Kuhn C., Boersch D., Glaetzer K.H., Schaefer U., Schaefer I., Schaefer U., Schaefer II., Schaefer U., Schaefer II., The state of four genes selectively expressed in the male germ lin of Drosophila melanogaster.";
Mech. Dev. 35:143-151(1991).
                                                                                                                                                                                                                                                                                   MEDLINE=22426069; PubMed=12537572;
Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E.,
Smith C.D., Tupy J.L., Whitfield B.J., Bayraktaroglu L., Berman B.P.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Staplecon M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
systematic review,";
Genome Biol. 3:RESEARCH0083.1-R
Genome Biol. 3:RESEARCH0083.1-R
-!- TISSUE SPECIFICITY: Testis.
-!- DEVELOPMENTAL STAGE: Primar
-!- DOMAIN: This protein is mos
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Drosophila melanogaster (Fruit fly).
Bukaryota; Metazoa; Arthropoda; Harapoda; Insecta; Pterygo
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                             Lewis S.E.;
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                                                                                                                                                                                                                  "Annotation
                                                                                                                                                                                                                           of the
                                                                                                                                     3: RESEARCH0083.1-RESEARCH0083.22 (2002)
                                                                                                                                                                                                              Drosophila melanogaster
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mostly composed of
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                 repetitive
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"Association of misexpression with steri
simulansand D. mauritiana.";

J. Mol. Evol. 59:277-282(2004).

EMBL; AY549957; AAS55566.1; -; Genomic_I
InterPro; IPR001450; 4Fe45 ferredoxin.
InterPro; IPR005634; MSSP.
InterPro; IPR001007; VWF C.
Pfam; PF03940; MSSP; 1.
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Q6QAT9;
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
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PROSITE; PS00198; 4FE45 FERREDOXIN; UNKNOWN_1.
PROSITE; PS01208; VWFC 1; 1.
PROSITE; PS01208; VWFC 1; 95A12F27D07B6F6C
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Drosophila simulans (Fruit fl); Hexapoda; Insecta; Pterygo
Bukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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Pfam; PF03940; MSSP; 1.
Developmental protein; Differentiation; Multigene family; Repeat;
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Ensembl; CG17945; Drosophila n
FlyBase; FBgn0004174; Mst84Dc.
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MEDLINE=20196006; PubMed=10731132; DOI=10.1126/science.287.5461.2185; Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F., George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Wan K.H., Doyle C., Baxter B.G., Helt G., Nelson C.R., Miklos G.L.G., Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P., Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MS87F DROME STANDARD; PRT; 56 AA.
PO8175; Q9VFQ6;
01-AUG-1988 (Rel. 08, Created)
01-AUG-1988 (Rel. 08, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Male specific sperm protein Mst67F.
Name=Mst87F; Synonyms=mst(3)gl-9; ORFNames=CG17956;
Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera, Endopterygota; Diptera, Enachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Michalak P., Noor M.A.F.;

"Association of misexpression with sterility in hybrids of Drosophila simulansand D. mauritiana.";

J. Mol. Evol. 59:277-282(2004).

EMBL; AY549955; AAS55564.1; -; Genomic_DNA.

InterPro; IPR001450; 4Fe45; ferredoxin.

InterPro; IPR001450; 4Fe45; ferredoxin.

InterPro; IPR001450; VWF_C.

Pfam; PF03940; MSSP; 1.
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NCBI TaxID=7226;
                                                                                                                                                                                                                                                                     Kuhn R., Schaefer U., Schaefer M.;
"Cis-acting regions sufficient for spermatocyte-specific transcriptional and spermatid-specific translational cont Drosophila melanogaster gene mst(3)gl-9.";
EMBO J. 7:447-454(1988).
                                                                                                                                                                                                                                                                                                                                           STRAIN=Canton-S;
MEDLINE=88211557; PubMed=2835228;
Kuhn R., Schaefer U., Schaefer M.
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PROSITE; PS01208; VWFC_1; 1.
SEQUENCE 55 AA; 5165 MW; 95A12F27D07B6F6C
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NCBI_TaxID=7227;
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RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Kimmel B.E., Kodira C.D., Kraft C., Krawitz S., Kulp D., Lai Z.,
Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Spier E., Spradling A.C., Stagleton M., Skupski M.P., Smith T.,
RA Spier E., Spradling A.C., Stagleton M., Skupski M.P., Smith T.,
RA Syirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissembach J.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissembach J.,
RA Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,
RA Gibbs R.A., Myers E.W., Rubin G.M., Venter J., Shu X., Smith H.O.,
RA Gibbs R.A., Myers E.W., Rubin G.M., Venter J., Shu X., Smith H.O.,
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RA Gibbs R.A., Myers E.W., Rubin G.M., Venter J., Shu X., Smith H.O.,
Ra Gibbs R.A., Weissenbach J., Shu R., Stone E., Spier E., Sprance of Drosophila melanogaster.";
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-!- TISSUE SPECIFICITY: Testis.
-!- DEVELOPMENTAL STACE: primary spermatocytes.
-!- DOMAIN: This protein is mostly composed of repe
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Misra S., Crosby M.A., Mungall C.J., Malthews B.B., Campbell K.S.
Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                  Developmental protein;
Spermatogenesis.
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AE003702; AAF54994.1; -; Genomic DNA
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A Lewis L.A., Gillaspy A.F., McLaughlin R.E., Gipson M., Ducey T.F.

A Ownbey T., Hartman K., Nydick C., Carson M.B., Vaughn J., Thomso

A Song L., Lin S., Yuan X., Najar F., Zhan M., Ren Q., Zhu H., Qi

A Kenton S.M., Lai H., White J.D., Clifton S., Roe B.A., Dyer D.W.

"The complete genome sequence of Neisseria gonorrhoeae.";

Submitted (MAR-2003) to the EMBL/GenBank/DDBJ databases.

EMBL; AB004969; AAW89584.1; -, Genomic_DNA.

GO; GO:0005489; F:electron transporte_activity; IEA.

GO; GO:0046972; F:metal ion binding; IEA.

R GO; GO:0046118; P:electron transport; IEA.

R GO; GO:006118; P:electron transport; IEA.

R InterPro; IPR001052; Rubredoxin.

R Pfam, PF00301; Rubredoxin.
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van Passel M.W.J., Bart A., Luyf A.C.M., van Kampen A.H.C.,
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SEQUENCE
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PRODOM; PD001610; RUBREDOXIN; 1.
PROSITE; PS00202; RUBREDOXIN_L
PROSITE; PS50903; RUBREDOXIN_L
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Neisseria gonorrhoeae (strain ATCC 700825 / FA 1090).
Bacteria, Proteobacteria; Betaproteobacteria; Neisser
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Bacteria; Proteobacteria;
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                                                                                                                                                                                                                                                                                                 PS00202; RUBREDOXIN; 1.
PS50903; RUBREDOXIN_LIKE; 1.
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No. 3e+02;
                                                                                                                                                                                                                                                          Iron; Metal-binding;
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                                                3e+02;
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                                                                                   Length 56;
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H., Qi S.,
/er D.W.;
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RESULT 16
Q9JQN3_NEIMA
Q9JQN3_NEIMA PRELIMINARY;
ID Q9JQN3;
AC Q9JQN3;
DT 01-OCT-2000 (TrEMBLrel. 15, L;
DT 01-OCT-2003 (TrEMBLrel. 25, L;
DT 01-OCT-2003 (TrEMBLrel. 25, L;
DE Putative rubredoxin.
GN OrderedLocusNames=NMA1201;
OS Neisseria meningitidis (serog:
OC Bacteria; Proteobacteria; Beta
OC Neisseriaceae; Neisseria.
OX NCBI_TaxID=65699;
                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 4
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                                              Neisseria meningitidis (serogroup A)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR004039; Rubredox.
InterPro; IPR001052; Rubredoxin.
Pfam; PF00301; Rubredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GO; GO:0005489; F:electron transporter activity; IEA. GO; GO:0046872; F:metal ion binding; IEA. GO; GO:0006118; P:electron transport; IEA.
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EMBL; AE002098; AAF41396.1;
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MEDLINE=20175755; PubMed=10710307; DOI=10.1126/science.287.5459.1809;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRINTS; PR00163; RUBREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Complete genome sequence of Neisseria meningitidis serogroup B strain
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                 Proteobacteria; Betaproteobacteria; Neisseriales;
ceae; Neisseria.
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Proteobacteria; Betaproteobacteria; Neisseriales;
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RUBREDOXIN_LIKE; 1.
; Electron transport; Iron; Metal-binding;
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Pred. No. 3e+
0; Mismatches
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Q6QAT8 DROSI
D Q6QAT8 DROSI PRELIMINARY;
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Best Local Similarity
Matches 4; Conserv
                                                                                                                                     STRAIN=Valparaiso;
STRAIN=Valparaiso;
STRAIN=Valparaiso;
PubMed=15486701; DOI=10.1007/s00239-004-2622-y;
Michalak P., Noor M.A.F.;
Michalak P., Noor M.A.F.;
"Association of misexpression with sterility in hybrids of Drosophila simulansand D. mauritiana.";
J. Mol. Evol. 59:277-282 (2004).
EMBL; AY549958; AAS55567.1; -; Genomic_DNA.
InterPro; IPR001450; 4Fe45 ferredoxin.
InterPro; IPR001007; VMPC_T; I.
PROSITE; PS00198; 4Fe45 FERREDOXIN; UNKNOWN_1.
PROSITE; PS01208; VMPC_T; 1.
SEQUENCE 58 AA; 5422 MW; 8C70EDCC7E69AC83 CRC64;
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05-JUL-2004
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"Complete DNA sequence of a serogroup A strain of Neisseria"
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NUCLEOTIDE SEQUENCE.
STRAIN=Z2491 / Serog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=Mst84Dc; ORFNames=CG17945;
Drosophila simulans (Fruit fly),
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Eukaryota; Metazoa; Arthropoda; Brachycera; Muscomorpha;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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HSSP; P24297; 1RWD.
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MEDLINE=20222556; PubMed=10761919; DOI=10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Complete proteome; Electron transport; Iron; Metal-binding; Rubredoxin.
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EMBL; AL162755; CAB84462.1; -;
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PROSITE; PS50903; RUBREDOXIN_LIKE; 1.
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ProDom; PD001610; Rubredox; 1.
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InterPro; IPR001052; Rubredoxin.
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Pred.
                                                                              Score 31; DB 2;
Pred. No. 3.1e+02
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                                                                                 3.1e+02;
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Query Match
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SEQUENCE
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Submitted
                                                                                                                                                                                                                EMBL; AF135821; AAG01185.1; -; mRNA.

EMBL; AF079059; AAD47373.1; -; Genomic_DNA.

EMBL; AF159976; AAA61823.1; -; mRNA.

EMBL; AF419252; AAA32699.1; -; Genomic_DNA.

HSSP; P15222; 1SIS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shunyi Z., Wenxin L.; "cDNA encoding a putative insect toxin from BmK."; Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Venom gland;
MEDLINE=20175769; PubMed=10708793; DOI=10.1016/S0041-0101(99)00212-3;
Zeng X.-C., Li W.-X., Zhu S.-Y., Peng F., Zhu Z.-H., Wu K.-L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mesobuthus martensii (Manchurian scorpion) (Buthus marte
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida;
Buthida; Buthoidea; Buthidae; Mesobuthus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Neurotoxin BmKCT precursor (C1-channel toxin)
chain toxin KCT) (BmKCL1) (TYCL1)
                                                                                                                                                                                                                                                                                                                                                  This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Elthe European Bioinformatics Institute. There are no rest
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Cloning and characterization of precursor of a chlorotoxin-like Buthus martensii Karsch.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=34649;
                                                                                                                                                                                  Pfam; PF05294; Toxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Venom
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5294; Toxin_5; 1.
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(JUL-1998)
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Neurotoxin BmKCT.
By similarity.
By similarity.
By similarity.
By similarity.
By similarity.
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Length
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RESULT 20
Q71DA0 DROYA
Q71DA0 DROYA PRELIMINARY;
ID Q71DA0;
AC Q71DA0;
DT 05-JUL-2004 (TrEMBLrel. 27)
DE CG4959 (Fragment).
CG4959 (Fragment).
CG Enkaryota; Metazoa; Arthro
OC Eukaryota; Endopterygota; I
OC Ephydroidea; Drosophilidae
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Matches 4; Conserv
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"A novel neurotoxin from scorpion BmK.";

Submitted (DEC-2000) to the EMBL/GenBank/DDBJ
-I- FUNCTION: Chloride channel ligand. Blocks
-chloride channels (By similarity).
-i- SUBCELLULAR LOCATION: Secreted.
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28-FEB-2003 (Rel. 41, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
Neurotoxin Bml2-b precursor.
  Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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between the Swiss Institute of Bioinformatics and the EMBL
the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
Xu C.-Q., Chi C.-W.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mesobuthus martensii (Manchurian scorpion) (Buthus martensii).
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Scorpiones;
Buthida; Buthoidea; Buthidae; Mesobuthus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q9BJW4;
                                                                     CG4959 (Fragment).
Drosophila yakuba (Fruit fly).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chloride channel inhibitor; Ionic channel inhibitor; Neurotoxin;
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5294; Toxin_5; 1.
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Neurotoxin Bm12-b.
By similarity.
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C2BAABAFCB88D9EA CRC64;
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Pred. No. 3.2
D; Mismatches
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RESULT 22

QBT403_DROME

QBT403;
DC QBT403;
DT 01-JUN-2002 (TYEMBLY). 21, Created)
DT 01-JUN-2002 (TYEMBLY). 21, Last sequence update)
DT 10-MAY-2005 (TYEMBLY). 30, Last annotation update)
DT 10-MAY-2015 (TYEMBLY).

QR AT235519 (CG31740-PA).

QR ORFNames=CG31740;
QS DYSOPHILA melanogaster (Fruit fly).

CE Ukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pte.
QC Neoptera; Endopterygota; Diptera; Brachycera; Muscomori,
QC Ephydroidea; Drosophilidae; Drosophila.

QX NCBI_TaxID=7227;
RM [1]
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ORFNames=CG4959;

Drosophila melanogaster (Fruit fly).

Drosophila melanogaster (Fruit fly).

Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;

Bukaryota; Endopterygota; Diptera; Brachycera; Muscomorpha;

Ephydroidea; Drosophilidae; Drosophila.
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FlyBase; FBgn0028883; CG4959.
SEQUENCE 60 AA; 6117 MW;
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MEDLINE=22887302; PubMed=1
Domazet-Loso T., Tautz D.;
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EMBL; AF531969; AAQ09868.1; -; mRNA
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RA Ballaw R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Botshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
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RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Cawley S., Fleischmann W.,
RA Glodek A., Gong F., Gorrell J.H., Glasser K.,
RA Harris M.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris M.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Liu X., Mattei B., McIntosh T.C., McLeodd M.P., McPherson D.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Mount S.M., My M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,
RA Melson D.R., Nelson K.A., Saunders R.D.C., Scheeler F., Shen H.,
RA Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
RA Shue B.C., Siden-Kiamos I., Stapleton M., Strong R., Sun S.,
RA Walliams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,
RA Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,
RA Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,
The Globes R. M., Wolley R. M., Shan M., Zhang G., Zhao
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                                                                                                                                                                                                                                                                                                      Celniker S.E., Wheeler D.A., Krommiller B., Carlson J.W., Halpern A Patel S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson A., George R.A., Hoskins R.A., Laverty T., Muzny D.M., Nelson C.R., Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E.J., Svirskas R., Tabor P.E., Wan K., Stapleton M., Sutton G.G., Venter Weinstock G., Scherer S.E., Myers E.W., Gibbs R.A., Rubin G.M.; "Finishing a whole-genome shotgun: release 3 of the Drosophila melanogaster euchromatic genome sequence."; Genome Biol. 3;RESEARCH0079-RESEARCH0079(2002).
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Patel S., Frise E., Wheel
Ashburner M., Celniker S
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  MEDLINE=22426069;
                                                                          transposable elements of the Drosophila meland nomics perspective."; me Biol. 3:RESEARCH0084.1-RESEARCH0084.20(2002)
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  PubMed=12537572
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P81761; Q8T0X4;
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Submitted
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Berkeley Drosophila Ge
Celniker S., Carlson J
Hoskins R., Stapleton
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lepidopteran-selective toxin precursor (ButaIT) (BTChl2).
Mesobuthus tamulus (Eastern Indian scorpion) (Buthus tamu
Eukaryota, Metazoa, Arthropoda, Chelicerata, Arachnida, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-2003 (Rel. 41, Created)
10-OCT-2003 (Rel. 42, Last sequence up
10-MAY-2005 (Rel. 47, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yu C., Rubin G.;
"Drosophila melanogaster release
Submitted (MAR-2000) to the EMBL/
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Newton K.A., Armugam
"Insect toxin from Me
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa;
Buthida; Buthoidea;
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                                                                                                                                                                                  "Isolation and toxin from the
                                                                                                                                                                                                                                            Wudayagiri R.,
                                                                                                                                                                                                                                                                      PubMed=11782289;
                                                                                                                                                                                                                                                                                       TISSUE=Venom;
                                                                                                                                                                                                                                                                                                                                                                         Submitted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUCLEOTIDE SEQUENCE.
                 MC Blochem. 2:16-16(2001).

1. FUNCTION: Induces flaccid paraly toxic to S.falculata larvae or m toxic to S.Falculata larvae or m subsectivity. Secreted.

1. SUBCELULAR LOCATION: Secreted.

1. TISSUB SPECIFICITY: Expressed by MASS SPECTROMETRY: MW=3856.7; ME NOTE=Ref.2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34
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                                                                                                                                                                                  characterization of a venom of South Indian
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Belongs
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on J., Wa
con M., P
                                                                                                                                                                                                                                                                                                                                                                                              Mesobuthus
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                                                                                                                        flaccid paralysis
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Wan K., Pfeiffer I
, Pacleb J., Park !
                                                                                                                                                                                                                                                                                                                                                                                              strong P.N.,
ithus tamulus
the
                                                                                                                                                                                                                                                                                                                                FUNCTION,
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                                                                                                                                                                                                                                                  Herrmann
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Pred. No. 3.2
); Mismatches
short
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                                      by the venom gland. METHOD=Electrospray;
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                                                                                                     mice
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red sc
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scorpion
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                                                                                                                                                                                  el lepidopteran-selective
scorpion, Mesobuthus
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                                                                                                                          H.virescens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arachnida; Scorpiones;
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Svirskas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CRC64;
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m
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                                                                                                                            larvae.
                                             RANGE=25-61;
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                                                                                                                                                                                                                                                                                                                                    TISSUE
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k S.E.
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Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
A Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,
A George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
A Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
B Erandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
A Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
A Barandon R.C., Rogers Y.-H.C., Andrews-Pfannkoch C., Baldwin D.
A Ballew R.M., Basu A., Baxerdale J., Bayraktaroglu L., Beasley E.M.,
B Ballew R.M., Basu A., Baxerdale J., Bayraktaroglu L., Beasley E.M.,
B Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
B Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
B Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
Cha Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                        STRAIN-Oregon-R;
MEDLINB-9210253; PubMed=1684716; DOI=10.1016/0925-4773(91)90064-D;
Kuhn R., Kuhn C., Boersch D., Glaetzer K.H., Schaefer U., Schaefer M.;
"A cluster of four genes selectively expressed in the male germ line of Drosophila melanogaster.";
Mech. Dev. 35:143-151(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JT 24

N DROME

MS84A DROME STANDARD; PRT; (01642; 09VIA3; 01-JUL-1993 (Rel. 26, Last sequence up. 10-MJV-2005 (Rel. 47, Last annotation 10-MAY-2005 (Rel. 26, Last annotation 10-MAY-2005)
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DISULFID
DISULFID
DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                       Male specific sperm protein Mst84Da.

Name=Mst84Da; ORFNames=CG17946;
Drosophila melanogaster (Fruit fly).

Eukaryota; Metazoa; Arrhropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                                                                        MEDLINE=20196006; PubMed=10731132; DOI=10.1126/science.287.5461.2185;
                                                                                                                                                                                                                     STRAIN=Berkeley;
                                                                                                                                                                                                                                                                                                                                                                              Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID=7227;
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                                                                                                                                                                                                                                        NUCLEOTIDE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pfam; PF05294; Toxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GO:0005576; C:extracellular region; GO:0019870; F:potassium channel inh GO:0009405; P:pathogenesis; IDA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    36
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                                                                                                                                                                                                                                     SEQUENCE
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                                                                                                                                                                                                                                     [LARGE SCALE GENOMIC DNA].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24
62
43
51
58
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By similarity.
By similarity.
By similarity.
By similarity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 31; DB 1;
Pred. No. 3.3e+02;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in no
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            63
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RESULT 25
Q8MI19 SHEEP
ID Q8MI19 SHEEP
AC Q8MI19;
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RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Liu X., Mattei B., McIncosh T.C., McLeod M.P., McPherson D.,
RA Liu X., Mattei B., McIncosh T.C., McLeod M.P., McPherson D.,
RA McTkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA McIson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Spier B., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Spier B., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Svirskas R., Tector C., Turner R., Venter B., Wang A.H., Wang X.,
Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Yeb J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,
RA Gibbs R.A., Myers B.W., Rubin G.M., Venter J.C.,
Fife genome sequence of Drosophila melanogaster.",
Fife genome sequence of Drosophila melanogaster.",
                                                                                                                                                                             Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                       Ensembl; CG17940; L., Mst84Da. FlyBase; FBgn0004172; Mst84Da. ThterPro; IPR005634; MSSP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=22426069; PubMed=12537572;
Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Misra S., Crosby M.A., Mungall C.J., Milburn G.H., Prochnik S.E.
Hradecky P., Huang Y., Kaminker J.S., Milburn G.H., Prochnik S.E.
Smith C.D., Tupy J.L., Whitfield B.J., Bayraktaroglu L., Berman B.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                               EMBL; X67703; CAA47937.1; -; EMBL; AE003672; AAF54022.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     systematic review."
                                                                                                                                                                                                                                                                                      Developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENOME REANNOTATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Annotation of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lewis S.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      s Swiss-Prot entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EN European Bioinformatics Institute. There are no resti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOMAIN: This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SIMILARITY: Belongs to the MST(3)CGP family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE SPECIFICITY: Testis.
DEVELOPMENTAL STAGE: Primar
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                SHEEP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     long
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                                                                                                                                                                                               Similarity
                                                                                                                                             2492
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ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     as its
                    PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3:RESEARCH0083.1-RESEARCH0083.22(2002)
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                                                                                                                                                                                                                                                    ₽,
                                                                                                                                                                                                                                                                                      protein; Differentiation; Multigene family;
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                                                                                                                                                                                                                                                    5806 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     content is in
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                                                                                                                                                                             Score 31; DE
Pred. No. 3.3
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                           melanogaster
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                                                                                                                                                                                                                                                                                                                                                                                             Genomic_DNA.
                  PRT;
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                                                                                                                                                                                                                Length 63;
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RESULT 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Placental artery endothelium; Chung J.-Y., Tsoi S.C.M., Wen Y.-X., Submitted (AUG-2002) to the EMBL/Geni EMBL; AF534638; AANO4109.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae; Caprinae; Ovis.
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01-OCT-2002 (TrEMBLrel. 22,
01-MAR-2004 (TrEMBLrel. 26,
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                                                                                                                                                                                                                                                                                                                                                               ProDom; PD001629; PD growth factor; 1. PROSITE; PS50278; PDGF 2; 1. Growth factor; Mitogen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE.
Tayade C., Black G.P., Croy
Submitted (MAY-2004) to the
EMBL; AY616676; AAT38876.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9823;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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GO:0008083; F:growth factor activity; IEA.
GO:0008283; F:cell proliferation; IEA.
GO:0000074; P:regulation of cell cycle; IEA.
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Pred. No. 3.5
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3.5e+02;
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RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Redeorge R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., RA Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D., RA Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D., RA Brandon R.C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., RA Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I., RA Gerry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., RA Dodson K., Doup L.B., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., RA Dodson K., Doup L.B., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., RA Gebler C., Gabrielian A.B., Garg N.S., Gelbart M.M., Glasser K., Golder C., Gabrielian A.B., Garg N.S., Gelbart M.M., Glasser K., Ra Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Alasin P., Kalush F., Karpen G.H., Ke Z., Kemison J.A., Ketchum K.A., Kimmel B.E., Kodira C.D., Kratt C., Kravitz S., Kulp D., Lai Z., Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D., RA Mount S.M., Woy M., Murphy B., Murphy L., Muzny D.M., Nelson D.L., RA Malazzolo M., Pitchan G.S., Pan S., Pollard J., Puri V., Reese M.G., RA Spier E., Spradling A.C., Stapleton M., Skupski M.P., Smith T., Spier E., Spradling A.C., Turner R., Venter E., Wang A.H., Wang X., Wang Z.-Y., Wassarman D.A., Weinstock G.M., Wissenbach J., Ra Glibbs R.A., Wyers E.M., Zhong W., Zhang G., Zhang G., Zheng L., Ra Gibbs R.A., Wyers E.W., Zhong W., Shand M., Sianng G., Zhao Q.A., Sheng L., Zhong S., Yao Q.A., Smith H.O., Ra Gibbs R.A., Wyers E.W., Zhong W., Shand M., Shang G., Zhao Q., Zheng L., Schence 287:2185-2195 (2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q9VM83;
01-MAY-2000 (TrEME
01-MAY-2000 (TrEME
10-MAY-2005 (TrEME
CG17376-PB, isofo
ORFNames=CG17376;
                                                                                                                                                                                                                                                                                                                                                                   science [2]
                                                                                                                                                                                                               Celniker S.E., Wheeler D.A., Kronmiller B., Carlson J.W., Halp Patel S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson C.R. George R.A., Hoskins R.A., Laverty T., Muzny D.M., Nelson C.R. Pacleb J.M., Park S., Ffeiffer B.D., Richards S., Sodergren E. Svirskas R., Tabor P.E., Wan K., Stapleton M., Sutton G.G., Ve Weinstock G., Scherer S.E., Myers E.W., Gibbs R.A., Rubin G.M.
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                               "Finishing a whole-genome shotgun: release 3 of the Drosophila melanogaster euchromatic genome sequence."; Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
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NCBI_TaxID=7227
                                                                 MEDLINE=22426070; PubMed=12537573;
Kaminker J.S., Bergman C.M., Kronmiller
Patel S., Frise E., Wheeler D.A., Lewis
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                          "The transposable elements of the Drosophila melanogaster
                                                  Ashburner M.,
                                                                                                                             NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                               MEDLINE=22426065; PubMed=12537568;
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               perspective.
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(TrEMBLrel. 13, Last sequence up)
(TrEMBLrel. 30, Last annotation
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3:RESEARCH0084.1-RESEARCH0084.20(2002)
                                                     Celniker S.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PubMed=10731132; DOI=10.1126/science.287.5461.2185;
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Hoskins R.A.,
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                                                                     update)
                                                                                      Carlson
                                                                       Rubin
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Matches 4
Query Match
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25-OCT-2004
25-OCT-2004
                                                                                                                                                                "Oryza sativa nipponbare(GA3) genc
Clone:OJNBa0016D04.";
                                                                                                                                                                                                                                                                                                        Oryza sativa (japonica cultivar-group). 
Eukaryota; Viridiplantae; Streptophyta; En 
Spermatophyta; Magnoliophyta; Liliopsida; 
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                        25-OCT-2004 (TrEMBLrel. 28, Created)
25-OCT-2004 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Hypothetical protein OSJNBa0016D04.29.
Name=CSJNBa0016D04.29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stapleton M., Čárlson J., Chavez C., Frise E., Park S., Wan K., Yu C., Rubin G.M., Celniker S. Submitted (FEB-204) to the EMBL/GenBank/DDBJ d EMBL; AR003615; AAF52440.1; -; Genomic DNA. EMBL; BT011549; AAS15685.1; -; mRNA. Ensembl; CG17376; Drosophila melanogaster. FlyBase; FBgn0042189; CG17376.

SEQUENCE 67 AA; 6871 MW; B9442F2C7B567C76 C
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Misra S., Crosby M.A., Mungall C.J., Milburn G.H., Prochnik S.E.
Hradecky P., Huang Y., Kaminker J.S., Milburn G.H., Prochnik S.E.
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                Hypothetical
SEQUENCE 6
                                                                                                                              Submitted (OCT-2002) to the EMBL; AP005800; BAD29328.1;
                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
Sasaki T., Matsumoto
                                                                                                                                                                                                                                                                               NCBI_TaxID=39947;
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                                                                                                           Gramene;
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Misra S., Crosby M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QGEPU8_ORYSA PRELIMINARY;
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4; Conser
                                                                                                           QGEPU8; -
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9 AA; 7109 MW;
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n M., Pacleb J., Park
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the EMBL/GenBank/DDBJ
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                                                                9FBA6730BE5730A6 CRC64;
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a; Poales; Poaceae;
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                    STRAIN=Oregon-R;
STRAIN=92102953; Pul
MEDLINE=92102953; Pul
Kuhn R., Kuhn C., Bo
"A cluster of four g
Kuhn R., Kuhn on R., Kuhn of Drosophila
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SEQUENCE
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Name=Mst84Dd; ORFNames=CG17935;

Drosophila melanogaster (Fruit fly).

Eukaryota; Metazoa; Arthropoda; Hexapoda; Inse
Neoptera; Endopterygota; Diptera; Brachycera;

Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ul-MAR-2004 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
ENSAMGP00000023827 (Fragment
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Q7PK85;
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The Anopheles gambiae Sequence Committee;
Submitted (APR-2004) to the EMBL/GenBank/DDBJ
-i- CAUTION: The sequence shown here is derive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The Anopheles gambiae Sequence Committee, "Anopheles gambiae re-annotation."; Submitted (APR-2002) to the EMBL/GenBank/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORFNames=ENSANGG00000020560;
                                                                                                        DEVELOPMENTAL
                                                                                                                                   NUCLEOTIDE
                                                                                                                                                                         NCBI_TaxID=7227;
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Anophelinae; Anopheles.
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953; PubMed=1684716; DOI=10.1016/0925-4773(91)90064-D; C., BoerEch D., Glaetzer K.H., Schaefer U., Schaefer four genes selectively expressed in the male germ lin melanogaster.";
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                                                                                                                                 [GENOMIC DNA],
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a; Nematocera; Culicoidea; Culicidae;
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35:143-151(1991)

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RA Amanatides P.G., Center S.E., Holt K.A., Evans C.A., Godyne J.J.,
RA Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Brandon R.C., Rogers Y.H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,
RA Ballew R.M., Basu A., Barman B.P., Bhandari D., Bolshakov S.,
Ra Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Gloden K.J., Evangelista C.C., Ferriar C., Ferriers S., Fleischmann W.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Gana P., Harris M.,
RA Harris N.L., Harvey D.A., Heiman T.J., Wei M.-H., Ibegwam C.,
RA Harris N.L., Harvey D.A., Heiman T.J., Wei M.-H., Ibegwam C.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Haris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Nelson D.H.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Spier B., Spradling A.C., Stapleton M., Stupski M.P., Smith T.,
RA Syirakas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
RA Wilsans R., Tector C., Turner R., Venter E., Wang A.H., Cang C.,
RA Wilsans R., Tector C., Turner R., Venter E., Wang A.H., Smith H.O.,
RA Glibbs R.A., Weers T.J., Simpson M., Skupski M.P., Smith T.,
RA Glibs R.A., Weers T.J., Singson M., Skupski M.P., Smith T.,
RA Jeng X., Yang S., Zang F.N., Zhang G., Zhao Q., Zheng L.,
RA Jeng X., Shan M., Zhang G., Zhao Q., Zheng L.,
RA Jeng X., Jeng J., Jeng J., Jeng J., Jeng L.,
RA Jeng X., Jeng J., Jeng J., Jeng J., Jeng J.,
RA Jeng J
                                                                                                                                                                                                                                                                                                                                                                                                                                            Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S., Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E., Smith C.D., Tupy J.L., Whitfield E.J., Baytaktaroglu Berman B.P., Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A., Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q., Stabeleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                     EMBL; X67703; CAA47940.1; -; EMBL; AE003672; AAF54026.1; PIR; S25775; S25775.
                                                                                                                                                                                        This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                      Genome Biol. 3:RESEARCH0083.1-RESEARCH0083.22(2002)
-!- TISSUE SPECIFICITY: Testis.
-!- DEVELOPMENTAL STAGE: Primary spermatocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=22426069; PubMe Misra S., Crosby M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENOME REANNOTATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=20196006; PubMed=10731132; DOI=10.1126/science.287.5461.2185;
                                                                                                                                                                                                                                                                                                  -! - DOMAIN:
                                                                                                                                                                                                                                                                                                                                                                                       systematic review
                                                                                                                                                                                                                                                        -!- SIMILARITY: Belongs to the MST(3)CGP family.
                                                                                                                                                                                                                                                                                                                                                                                                      "Annotation of the Drosophila melanogaster euchromatic genome: a
                                                                                                                                                                                                                                                                                                                                                                                                                                 Lewis S.E.;
                                                                                                                                                  as
                                                                                                                                                                    European
CG17935; Drosophila melanogaster FBgn0004175; Mst84Dd.
                      CG17935;
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                                                                                                                                                                        Bioinformatics
                                                                                                                                                  its content
                                                                                                                                                                                                                                                                                                protein
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                                                                                                                                                                                                                                                                                                Primary spermatocytes. is mostly composed of
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is in no wa
                                                              -; Genomic_DNA
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                                                                                    DNA.
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Q5TSU4_ANOGA PRELIMINARY;
Q5TSU4;
Q5TSU4;
Q5TSU4;
01-FEB-2005 (TremBLrel. 29)
01-FEB-2005 (TremBLrel. 29)
01-FEB-2005 (TremBLrel. 29)
                                                                                    13-SEP-2005
13-SEP-2005
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ORFNames=ENSANGG00000023294;
Anopheles gambiae str. PEST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR005634; MSSP.
Pfam; PF03940; MSSP; 1.
Developmental protein; Differentiation; Multigene family; Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      88
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Drosophila
Eukaryota;
                                                        Name=Mst84Dd;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (APR-2004) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
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                                                                                                                                                                                                                                                                       32
DROME
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melanogaster (Fruit fly).
Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
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Pred. No. 3.7e+0;
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Pred. No. 3.7e+02;
                                                                                                                                                                                                                                           PRT;
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gion; IEA.
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ID Q74DK8 AC 
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
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Q4KKY5;
13-SEP-2005 (7
13-SEP-2005 (7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE.

STRAIN=PCA / ATCC 51573;

PubMed=14671304; DOI=10.1126/science.1088727;

Methe B.A., Nelson K.E., Eisen J.A., Paulsen I.T., Nelson W.C.,

Meduberg J.F., Wu D., Wu M., Ward N.L., Beanan M.J., Dodson R.J.

Madupu R., Brinkac L.M., Daugherty S.C., Deboy R.T., Durkin A.S.,

Gwinn M.L., Kolonay J.F., Sullivan S.A., Haft D.H., Selengut J.,

Davidsen T.M., Zafar N., White O., Tran B., Romero C., Forberger H

Weidman J.F., Khouri H.M., Feldblyum T.V., Utterback T.R.,

Van Aken S.E., Lovley D.R., Fraser C.M.;

"Genome of Geobacter sulfurreducens: metal reduction in subsurface
      Mus musculus (Mouse)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stapleton M., Carlson J., Chavez C., Frise E., George R. Park S., Wan K., Yu C., Celniker S.; Submitted (MAY-2005) to the EMBL/GenBank/DDBJ databases. EMBL; BT023344; AAY55760.1; -; mRNA.
SEQUENCE 72 AA; 6840 MW; 8CE37449E5F88ABA CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Complete proteome; Hypothetical protein. SEQUENCE 72 AA; 8191 MW; F2E9B550E9D43BB9 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q74DK8;
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Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID=7227;
                                                                                                         Name=Pkd1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Science
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            environments.";
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                                                                                                                                    Hypothetical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Geobacteraceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OrderedLocusNames=GSU1308;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GSU1308;
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AE017180; AAR34684.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGPC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGPC 5
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                                                                                                                                                                                                                            (TrEMBLrel.
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                                                                                                                                    protein.
                                                                                                                                                                                                                                                                                               PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41
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Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Glires; Rodentia; Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
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                                                                                                                                                               Created)
Last sequence
Last annotation
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Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 31; DB 2; Pred. No. 3.7e+02; Mismatches 0;
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Pred. No.
                                                                                                                                                                                                                                                                                               PRT;
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annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 72;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pacleb J.,
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RESULT 35
Q6EPU3_ORYSA PRELIMINARY;
ID Q6EPU3;
AC Q6EPU3;
DT 25-OCT-2004 (TrEMBLrel. 2)
DT 25-OCT-2004 (TrEMBLrel. 2)
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RP TISSUB-Jaw and Limb;

RX MEDLINE-22388257; PubMed-12477932; DOI=10.1073/pnas.242603899;

RX MEDLINE-22388257; PubMed-12477932; DOI=10.1073/pnas.242603899;

RX Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA HOpkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Ugdnin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Blosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Hilalon D.K., Muzny D.M., Sodergren B.J., Lu X., Glbbs R.A.,

RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Glbbs R.A.,

RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Halkesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

"Generation and initial analysis of more than 15,000 full-length human

"Generation and initial analysis of more than 15,000 full-length human

"Generation and initial analysis of more than 15,000 full-length human
Query Match
Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 4; Conserv
                                                                                                                                                                                                             clone:OSJNBa0016D04.";
Submitted (OCT-2002) to the
EMBL; AP005800; BAD29333.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypothetical SEQUENCE 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Name=OSJNBa0016D04.34;
Oryza sativa (japonica cultivar-group).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-OCT-2004 (TrEMBLrel. 28, Created)
25-OCT-2004 (TrEMBLrel. 28, Last sequence up
25-OCT-2004 (TrEMBLrel. 28, Last annotation
Hypothetical protein OSJNBa0016D04.34.
                                                                                                                                                                                   Gramene;
                                                                                                                                                                                                                                                                                                                                       Sasaki T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; BC099589; AAH99589.1; MGI; MGI:97603; Pkd1.
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TISSUE=Jaw and Limb;
NIH MGC Project;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Muroidea; Muridae; Murinae;
NCBI TaxID=10090;
[1] ~.
                                                                                                                                                                                                                                                                                                    Sasaki T., Matsumoto T., Katayose Y.;
"Oryza sativa nipponbare(GA3) genomic
                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=39947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (JUL-2005) to the
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Similarity 4; Conserv
                                                                                                                                                                                   Q6EPU3; -
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100.0%; ilarity 100.0%; Conservative 0;
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                                                                                                                   protein.
3 AA; 7746 MW;
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                                                                                                                                                                                                          EMBL/GenBank/DDBJ
-; Genomic_DNA.
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                                                                                                                          3EF220FB4F36D10A CRC64;
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Score 31; DB 2;
Pred. No. 3.8e+02;
Mismatches 0;
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                                                         Length 73;
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NUCLEOTIDE SEQUENCE.

MEDLINE=85122780; PubMed=2578818; DOI=10.1016/0167-4781(85)90049-1; Wilton S.D., Crocker L.A., Rogers G.E.;

"Isolation and characterisation of keratin mRNA from the scale epidermis of the embryonic chick.";

Biochim. Biophys. Acta 824:201-208(1985).

EMBL; M25641; AAA48932.1; -; mRNA.

PIR; A60136; A60136.

Ensembl; ENSGALG0000009189; Gallus gallus.

GO; GO:0005200; F:structural constituent of cytoskeleton; IEA.

InterProj. IPR093461; Keratin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MS94B DROME STANDARD; PRT; /4 Am. Q01643; Q9MZE2; Q9VLA1; Q1-1993 (Rel. 26, Created) 13-SEP-2005 (Rel. 48, Last sequence update) 13-SEP-2005 (Rel. 48, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                          Male specific sperm protein Mst84Db.
Name-Mst84Db; ORFNames-CG17934;
Drosophila melanogater (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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Eukaryota; Metazoa; Choruacu,
Eukaryota; Aves; Neognathae;
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Gallus gallus (Chicken)
                                                                                                    MEDLINE=92102953; PubMed=1684716; DOI=10.1016/0925-4773(91)90064-D; Kuhn R., Kuhn C., Boersch D., Glaetzer K.H., Schaefer U., Schaefer M "A cluster of four genes selectively expressed in the male germ line of Drosophila melanogaster.";
Mech. Dev. 35:143-151(1991).
                                                                                                                                                                                                        STRAIN=Oregon-R;
MEDLINE=92102953;
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                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE [GENOMIC DNA], TISSUE SPECIFICITY, DEVELOPMENTAL STAGE.
                             STRAIN=Berkeley
                                                        NUCLEOTIDE
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                                                      SEQUENCE [LARGE SCALE GENOMIC DNA].
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PubMed=10731132; DOI=10.1126/science.287.5461.2185;
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Neognathae; Galliformes; Phasianidae; Phasianinae;
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Pred. No. 3.8e+02;
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RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,
RA George R.A., Levis S.E., Richards S., Ashburner M., Henderson S.N.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Barlaw R.M., Beas R.C., Rogers Y.H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Abril J.F., Agbayani A., An H.-J., Andrews-Ffannkoch C., Baldwin D.,
RA Ballew R.M., Beau A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Beson K.Y., Benos P.V., Berman B.P., Bhandari D., Bothshakov S.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Glerry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Gana P., Harris M.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
Harvey D.A., Heiman T.J., Wei M.-H., Ibegwam C.,
RA Harris N.L., Harvey D.A., Heiman T.J., Wei M.-H., Ibegwam C.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Merkilov G., Mishina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K., Maxon K., Nussekern D.R., Paleb J.M.,
RA Melson D.R., Nelson K., Saunders R.D.C., Scheeler F., Shen H.,
RA Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
RA Shue B.C., Stan-Kiamos I., Simpson M., Skupski M.P., Smith T.,
RA Sheng X.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Sings R.H., Zhong F.N., Zhong W., Zhang G., Zhao Q., Zheng L.,
RA Jahan M., Kalush F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,
RA Jahan M., Schoeler F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,
RA Jahan M., Shuba R.A., Shih H.O., Zhang G., Zhao Q., Zheng L.,
Ra Jahan M., Shih H., Jahan M., Zhang G., Zhao Q., Zheng L.,
Ra Jahan M., Shih H.
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Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Hradecky P., Huang Y., Kaminker J.S., Milburn G.H., Prochnik S.E.
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktaroglu L., Berman B.
Bettencourt B.R., Celniker S.E., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                                                                                                  Stapleton M., Carlson J.W., Brokstein P., Yu C., C George R.A., Guarin H., Kronmiller B., Pacleb J.M. Rubin G.M., Celniker S.E.; Rubin G.M., Celniker S.E.; Genome Biol. 3:RESEARCH0080.1-RESEARCH0080.8(2002)
                                                                                                                                                                                                                                                                                                                                                                                STRAIN=Berkeley;
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This the between European as long a Swiss-Prot entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the Eleuropean Bioinformatics Institute. There are no rest as content is 11 no way modified and this statement restrictions EMBL ρı collaboration outstation 18

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Proc. Natl. Acad. Sci. U.S.A. 101:14925-14930(2004).

REMBL; APO06618; BAD57292.1; -; Genomic_DNA.

RGO; GO:0005489; F:electron transporter activity; IEA.
RGO; GO:0005106; F:iron ion binding; IEA.
RGO; GO:0006118; p:electron transport; IEA.
RGO; GO:0006118; p:electron transport; IEA.
R InterPro, IPR001450; 4Fe45 Fe_S_bd.
R InterPro, IPR001450; 4Fe45 Fe_S_bd.
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EMBL; AE03672; AAF54024.1; -; Genomic_DNA.
EMBL; AY113221; AAM29226.1; ALT_SEQ; mRNA.
PIR; S25773; S25773.
HSSP; Q9S7B3; 1EHH.
Ensembl; CG17934; Drosophila melanogaster.
FlyBase; FBgn0004173; MSC8D.
InterPro; IPR005634; MSSP; 1.
Pfam; PF03940; MSSP; 1.
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COMPBIAS 2
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STRAIN-IFM 10152;
PubMed=15466710; DOI=10.1073/pnas.0406410101;
Ishikawa J., Yamashita A., Mikami Y., Hoshino Shiba T., Hattori M.;
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                                                                                                                                                                                                                       Score 31; DE
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C -> W (in Ref. 1).
P -> L (in Ref. 1).
CC64F0F7A7E7CABC CRC64;
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AC Q4RP51;
DT 13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
DT 13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
DT 13-S
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Best Local :
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Shinose M., Takahashi Y., H
Kikuchi H., Shiba T., Sakak
"Genome sequence of an indu
avermitilis: deducing the a
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PROSITE; PS00198; 4FE4S FEREDOXIN;
4Fe-48; Complete proteome; Electron
Metal-binding; Transport.
SEQUENCE 75 AA; 8139 MW; 8070930
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STRAIN-MA-4680 / ATCC 31267 / NCIMB 12804 / NRRL 8165;

MEDLINE-22608306; PubMed=12692562; DOI=10.1038/nbt820;

Ikeda H., Ishikawa J., Hanamoto A., Shinose M., Kikuch Sakaki Y., Hattori M., Omura S.;
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01-JUN-2003 (TrEMBLrel. 24, Last sequ
01-MAR-2004 (TrEMBLrel. 26, Last anno
putative ferredoxin.
Name=fdxE; OrderedLocusNames=SAV4856;
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GO; GO:0005489; F:electron transporter activity;
GO; GO:0005506; F:iron ion binding; IEA.
GO; GO:0006572; F:metal ion binding; IEA.
GO; GO:0046872; F:metal ion binding; IEA.
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR001450; 4Fe4S Fe_S_bd.
Pfam; PF00037; Fer4; 2.
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STRAIN=MA-4680 / ATCC 31267 / NCIMB 12804 / NRgL 8165;
MEDLINE=21477403; PubMed=11572948; DOI=10.1073/pnas.211433198;
Omura S., Ikeda H., Ishikawa J., Hanamoto A., Takahashi C.,
Omura S., Takahashi Y., Horikawa H., Nakazawa H., Osonoe T.,
Shinose M., Takahashi Y., Horikawa H., Nakazawa H., Osonoe T.,
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BA000030; BAC72568.1; -
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/cetaceae; Streptomyces.
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Pred. No. 3.9
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RESULT
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DT OCCUPAGE
   RESULT 42
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Q74114 MYCPA PRELIMINARY;
ID Q74114;
AC Q74114;
DT 05-JUL-2004 (TrEMBLrel 27,
DT 05-JUL-2004 (TREMBLRel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Summer E.J., Peek M.L., Haliburton J.R., Hall E., Heusinkveld Simser J., No E.G., Gonzalez C.F., Young R.F.; Submitted (JUL-2003) to the EMBL/GenBank/DDBJ databases. EMBL; AY357582; AAQ63363.1; -; Genomic DNA. GO; GO:0019067; P:viral assembly, maturation, egress, and rel.
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
Putative head-tail joining protein.
Burkholderia cepacia phage BccpNazgul.
Viruses; dsDNA viruses, no RNA stage; Caudovirales
                                                  Mycobacterium paratuberculosis.
Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
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ProDom; PD019366; gpW; 1.
SEQUENCE 76 AA; 8345 MW; 9DBFDD73E5A96799
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               Mycobacteriaceae;
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ID Q4SJ47_TETNG
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QBXHP3;
QBXHP3;
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QBXHP3;
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=1502;
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flesh-eater";
Proc. Natl. Acad. Sci. U.S.A. 99:996-1001(2002).
EMBL; BA000016; BAB82145.1; -; Genomic DNA.
GO; GO:0005489; F:electron transporter activity;
GO; GO:0006118; P:electron transport; IEA.
                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

STRALIN-13 / Type A;

MEDLINE-21664373; PubMed-11792842; DOI=10.1073/pnas.022493799;

Shimizu T., Ohtani K., Hirakawa H., Ohshima K., Yamashita A.,

Shiba T., Ogasawara N., Hattori M., Kuhara S., Hayashi H.;

"Complete genome sequence of Clostridium perfringens, an anaer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae; Clostridium.
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PROSITE; PS00198; 4FE4S FERREDOXIN;
4Fe-4S; Complete proteome; Electron
Metal-binding; Transport.
SEQUENCE 76 AA; 8289 MW; 8CFSD78
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                                                                                                                        InterPro; IPR011911; GlrX_YruB.
InterPro; IPR006663; Thioredox_dom2
TIGREAMB; TIGR02196; GlrX_YruB; 1.
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EMBL/GenBank/DDBJ
-; Genomic_DNA.
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Last annotation update)
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                                                                                    3337E0EB2A301508 CRC64;
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Score 31; DB
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    transport; Iron; Iron-sulfur;

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                    .9e+02;
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Jaillon O., Aury J.M., Brunet F., Petit J.L., Stange-Thomann N.,
Mauceli E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,
A Mauceli E., Bouneau L., Fischer S., Lutfalla G., Dossat C., Segurens B.,
A Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,
A Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
A Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
A Biemont C., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,
Cruaud C., Duprat S., Brottier P., Coutanceau J.P., Gouzy J.,
A Cruaud C., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
A Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
A Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
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A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Lindblad-Toh K., Birren B., Nusbaum C., Kahn D., Robinson-Re
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Q9VJM5;
01-MAY-2000
01-MAY-2000
01-MAR-2004
Wan K.H., Doy
Abril J.F., J
Ballew R.M.,
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13-SEP-2005
13-SEP-2005
13-SEP-2005
                      NUCLEOTIDE SEQUENCE.

MEDLING-20196006; pubMed=10731132; DOI=10.1126/science.287.5461.2185; Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., Adams M.D., Celniker S.E., Li P.W., Hoskins R.A., Galle R.P., Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Henderson S.N., George R.A., Lewis S.E., Richards S., Ashburner M., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D., Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., Wan K.H., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Andrews-Pfannkoch C., Baldwin D., Champe M., Proposition of the second of the secon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Name=CG13245; ORFNames=CG13245;
Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygo:
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
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13-SEP-2005 (TrEMBLrel. 31, Last sequence update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
13-SEP-2005 (TrEMBLrel. 31, Last annotation update)
Chromosome undetermined SCAF14576, whole genome sho
ORFNames=GSTENG00017363001;
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Bukaryota; Metazoa; Chordata; Craniata; Verte
Actinopterygii; Neopterygii; Teleostei; Eutel
Acanthomorpha; Acanthopterygii; Percomorpha;
Tetradontoidea; Tetraodontidae; Tetraodon.
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L-MAY-2000 (TrEMBLrel.
MAR-2004 (TrEMBLrel.
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L; CAAE01014576; CAF99335
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JESCOPE, Whitehead Institute Centre for Genome Research, itted (FEB-2004) to the EMBL/GenBank/DDBJ databases.

CAUTION: The sequence shown here is derived from an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25
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9043 MW; 25F48C1751A7D3A3 CRC64;
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Last sequence update)
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Pred. No.
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Euteleostei; Neoteleostei;
rpha; Tetraodontiformes;
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           Beasley E.M.
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RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Fleischmann W.,
RA Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Gelbart W.M., Glasser K.,
RA Harris N.L., Harvey D.A., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Lisko P., Lei Y., Levitsky A.A., Li J.H., Li Z., Liang Y., Lin X.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Nelson D.L.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Spier E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Svirekas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Wang S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,
RA Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C.;
The genome sequence of Drosophila melanogaster.";
RN [71]
                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE:
MEDLINE=22426069; PubMed=12537572;
MEDLINE=22426069; PubMed=12537572;
Mista S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S.,
Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E.,
Smith C.D., Tupy J.L., Whitfield E.J., Bayraktearoglu L., Berman B.P.,
Bettencourt B.R., Celniker S.B., de Grey A.D.N.J., Drysdale R.A.,
Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.Q.,
Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
"Drosophila melanogaster release Submitted (MAR-2000) to the EMBL/[6]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Celniker S.E., Wheeler D.A., Kronmiller B., Carlson J.W., Halpern A., Patel S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson A., George R.A., Hoskins R.A., Laverty T., Muzny D.M., Nelson C.R., Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E.J., Svirskas R., Tabor P.E., Wan K., Stapleton M., Sutton G.G., Venter C., Weinstock G., Scherer S.E., Myers E.W., Gibbs R.A., Rubin G.M.; "Finishing a whole-genome shotgun: release 3 of the Drosophila melanogaster euchromatic genome sequence."; Genome Biol. 3:RESEARCH0079-RESEARCH0079(2002).
                                                                        Berkeley Drosophila Genome Project;
Celniker S., Carlson J., Wan K., Pfeiffer
Hoskins R., Stapleton M., Pacleb J., Park
                                                                                                                                                                                         "Annotation of the systematic review."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patel S., Fris
Ashburner M.,
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M
                                                      Rubin
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2426065; PubMed=12537568;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    J.S., Bergman C.M., Kroni
Frise E., Wheeler D.A.,
M., Celniker S.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                           perspective.";
1. 3:RESEARCH0084.1-RESEARCH0084.20(2002)
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                                                                                                                                     SEQUENCE
                                                                                                                                                                           3:RESEARCH0083.1-RESEARCH0083
                                                                                                                                                                                                             Drosophila melanogaster
                      EMBL/GenBank/DDBJ
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                                        4 sequence
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                                                                          S.B
                                                                                                                                                                                                             euchromatic
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., Rubin
                                                                            Frise E.,
Svirskas
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Q95JG5 BOVIN
ID Q95JG5;
AC Q95JG5;
DT 01-DEC-2
DT 01-OCT-2
DE Natural
GN Name=KUR
GO Bos taur
OC Pecora;
GN NCBI_Tax
NN [1]
TAX
RN [1]
TAX
RN [1]
RP NUCLEOTI
RC TISSUE-S
RX MEDLINE-
RA Govaerts
RA Govaerts
RA Govaerts
RA Govaerts
RT "Homolog
RT expresse
RT Vet. Imm
DR EMBL; AF
DR SMR; Q95
DR GO; GO; OD
DR FISSP; OS
DR SMR; P95
DR SMR; P95
DR SMRT; P95
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DR PFAM; PF
DR PROSITE;
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095JG6 BOVIN
ID 095JG6;
AC 095JG6;
DT 01-DEC-2001 (TrEMBLrel. 19,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expressed in cattle.";

Vet. Immunol. Immunopathol. 80:339-344 (2001).

EMBL; AF322885; AAK73810.1; -; mRNA.

HSSP; O54709; 1HQ8.

SMR; Q95JG5; 1-80.
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FlyBase;
Submitted (MAR-2005) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chorda
Mammalia; Eutheria; Lauraa
Pecora; Bovidae; Bovinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE=Spleen;
MEDLINE=21351526; PubMed=11457486; DOI=10.1016/S0165-2427(01)00295-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Natural killer cell
                                                                                                                                                                                                                                                                                                                                                      PROSITE;
                                                                                                                                                                                                                                                                                                                                                                         SMART; SM00034; CLECT;
                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00059; Lectin_C; 1
                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR001304; Lectin_C.
                                                                                                                                                                                                                                                                                                                                                                                                                                              GO; GO:0004872; F:receptor activity; GO; GO:0005529; F:sugar binding; IEA. InterPro; IPR002353; AntifreezeII.
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PS50041; C_TYPE_LECTIN_2;
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Pred. No. 4.1e+02;
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Best Local S
Matches 4
                                                                                      Eukaryota; Metazoa; C
Mammalia; Eutheria; I
Pecora; Bovidae; Bovi
NCBI TaxID=9913;
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NON_TER
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SEQUENCE
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Testicular luteinizing hormone beta subunit.
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01-OCT-2003 (TrEMBLrel. 25, 1
Natural killer cell receptor
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                           InterPro; IPR006208; Cys_knot.
InterPro; IPR001545; Gly_hormc
Pfam; PF00007; Cys_knot; 1.
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9429
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Laurasiatheria;
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Pred. No. 4.1
0; Mismatches
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046622_CERSI
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A Mauceli E., Bouneau L., Fischer C., Ozouf-Costaz C., Bernot A.,
A Nicaud S., Jaffe D., Fischer S., Lutfalla G., Dossat C., Segurens B.,
A Dasilva C., Salanoubat M., Levy M., Boudet N., Castellano S.,
A Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,
A Biemont C., Skalli Z., Cattolico L., Poulain J., De Berardinis V.,
A Cruaud C., Duprat S., Brottier P., Coutanceau J.P., Gouzy J.,
A Cruaud C., Duprat S., Brottier P., Cottanceau J.P., Gouzy J.,
A Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
A Parra G., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,
A Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,
A Lindblad-Tok K., Bürren B., Nusbaum C., Kahn D., Robinson-Rechavi M.,
A Laudet V., Schachter V., Quetier F., Saurin W., Scarpelli C.,
Wincker P., Lander E.S., Weissenbach J., Roest Crollius H.;
"Genome duplication in the teleost fish Tetraodon nigroviridis reveals
The early vertebrate proto-karyotype.";
Nature 431:946-957(2004).
                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 4; Conser
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Best Local
       01-JUN-1998
01-JUN-1998
01-JUN-2003
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B7_TETNG
Q4T7B7_1
Q4T7B7;
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046622;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleosteti; Buteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
Tetradontoidea; Tetraodontidae; Tetraodon.
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13-SEP-2005
13-SEP-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genoscope; Whitehead Institute Centre for Genome Research; Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases.
-i- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry whi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (Fragment).
ORFNames=GSTENG00005813001;
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; CAAE01008161; CAF91215.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CAUTION: The sequence shown EMBL/GenBank/DDBJ whole genc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        49
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4; Conserv
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AA;
       (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                        PRELIMINARY;
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8515 MW; F9EA66C2FD6FC97D CRC64;
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8611 MW;
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Pred. No. 4.1
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Pred. No.
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Q4ZC47 9VI
Q4ZC47 9VI
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DT 13-5
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DT 13-7
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060744 HUMAN

ID 060744;
AC 060744;
AC 060744;
DT 01-AUG-1998 (TrEMBLrel. 07, C1)
DT 01-AWG-1998 (TrEMBLrel. 26, L4)
DT 01-MAR-2004 (TrEMBLrel. 26, L4)
DE Thioredoxin delta 3 (Fragment)
GN Name=TXN delta 3;
OS Homo sapiens (Human).

CC Eukaryota; Metazoa; Chordata;
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13-SEP-2005
13-SEP-2005
13-SEP-2005
                                                                            Thioredoxin delta 3 (Fragment).
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Proc. Natl. Acad. Sci. U.S.A. 102:5174-5179(2005).
EMBL, AY94959; AAX91390.1; -; Genomic DNA.
SEQUENCE 83 AA; 8546 MW; 710C2C35BEAB2C31 CRC6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PubMed=15788529; DOI=10.1073/pnas.0501140102; Kwan T., Liu J., Dubow M., Gros P., Pelletter J.; Withe complete genomes and proteomes of 27 Staphylococcus aureus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROSITE; PS00689; GLYCO_HORMONE_BETA_2;
NON_TER 1 1
SEQUENCE 82 AA; 8643 MW; 270411BA19
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Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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InterPro; IPR001545; Gly_hormoneB.
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to the EMBL/GenBank/DDBJ databases.
365.1; -; Genomic_DNA.
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Pred. No. 4.2e-
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QÉMWEB ORYSA PRELIMINARY;
QÉMWEB;
05-JUL-2004 (TrEMBLrel. 2
05-JUL-2004 (TrEMBLrel. 2
05-JUL-2004 (TrEMBLrel. 2
05-JUL-2004 (TrEMBLrel. 2
Name=B1358B12.4;
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SEQUENCE
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GO; GO:0005489; F:electron transporter activity; IEA.
GO; GO:000518; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin, 1.

PRINTS; PR00421; THIOREDOXIN.
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SMR; C
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PubMed=15215690; DOI=10.1023/B:VIRU.0000032795.17713.6c;
PubMed=15215690; DOI=10.1023/B:VIRU.0000032795.17713.6c;
Varling M., Bartkowiak K., Pape H., Meinhardt F.;
"The Genome of phiAsp2, an Actinoplanes Infecting Phage.";
Virus Genes 29:117-129(2004).
EMBL; AY576796; AAT367581; -; Genomic DNA.
SEQUENCE 84 AA; 8995 MW; C1ECF1218IEB7C72 CRC64;
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"An alternative splice variant of human
Chin. Sci. Bull. 43:292-295 (1998).
EMBL; AF065241; AAC17430.1; -; mRNA.
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05-JUL-2004
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Q6J821;
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100.0%; P
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Last sequence update)
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Pred. No. 4.2e+02;
; Mismatches 0;
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Pred. No.
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annotation update)
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O QT6275;

C Q16275;

OTENBELTEL. 01, Created)

OT 01-NOV-1996 (TREMBLrel. 01, Last sequence updat

DT 01-NAR-2004 (TREMBLrel. 26, Last annotation upd

DE Type I hair keratin (Fragment).

GN Name=type I hair keratin;

OS Homo sapiens (Human).

OS Homo sapiens (Human).
  SORRE RAPARA RAP
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Best Local S
Matches 4
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Gramene; Q6MME8; -.
InterPro; IPR006461; DUF_A_thal_Cys.
Pfam; PF04749; PLAC8; 1.
Pfam; PF04749; A4 AA; 9417 MW; F0BA5C9
                          Bowden P.E., Hainey S., Parker G., Hodgins M

"Sequence and expression of human hair kerat

J. Dermatol. Sci. 7:0-0(0).

-i- SIMILARITY: Belongs to the intermediate

EMBL; S75797, AAB32812.1; -; Genomic DNA.

GO; GO:0005802; C:intermediate filament; IEA

GO; GO:0005198; F:structural molecule activi

InterPro; IPR001664; IF.
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Liu Y.L., Mu J., Yu Z., Chen L., Fan D.L., Weng Q.J., Zhang L.,
Lu Y.Q., Yu S.L., Liu X.H., Lu T.T., Zhang Y.J., Lu Y., Li C., Li T.
Zhang Y., Hu H., Jia P.X., Qian Y.M., Ying K., Zhou B., Chen Z.H.,
Hao P., Zhang L., Wu M., Zhang R.Q., Guan J.P., Fu G., Wang S.Y.,
Ren S.X., Lu G., Lin W., Gu W.Q., Zhu G.F., Tu Y.F., Jia J., Yin H.!
Zhang Y., Cai Z., Chen J., Kang H., Chen X.Y., Shao C.Y., Sun Y.,
Hu Q.P., Zhang X.L., Zhang W., Wang L.J., Ding C.W., Sheng H.H.,
Gu J.L., Chen S.T., Ni L., Zhu F.H., Hong G.F.,
Submitted (NOV-2003) to the EMBL/GenBank/DDBJ databases.
EMBL, EX842605; CAE75995.1; -; Genomic_DNA.
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NÜCLEOTIDE SEQUENCE:

PubMed=12447439; DOI=10.1038/nature01183;

Peng Q. Zhang Y., Hao P., Wang S., Fu G., Huang Y., Li Y., Zhu J.,

Liu Y., Hu X., Jia P., Zhang Y., Zhao Q., Ying K., Yu S., Tang Y.,

Weng Q., Zhang L., Lu Y., M. J., Lu Y., Zhang L.S., Yu Z., Fan D.,

Liu X., Lu T., Li C., Wu Y., Sun T., Lei H., Li T., Hu H., Guan J.,

Wu M., Zhang R., Zhou B., Chen Z., Chen L., Jin Z., Wang R., Yin H.,

Cai Z., Ren S., Lv G., Gu W., Zhu G., Tu Y., Jia J., Zhang Y.,

Chen J., Kang H., Chen X., Shao C., Sun Y., Hu Q., Zhang X., Zhang W.,

Mang L., Ding C., Sheng H., Gu J., Chen S., Ni L., Zhu F., Chen W.,

Lan L., Lai Y., Cheng Z., Gu M., Jiang J., Li J., Hong G., Xue Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lan L.,
Han B.;
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
MEDLINE=95092656; PubMed=7528047;
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Pred. No.
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nair keratin
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rimates; Catarrhini; Homin
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Gaps

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Hominidae,

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RESULT
Q8W710
ID Q8W710
AC Q8
DT Q1
DT Q1
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RP RP TRX
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Name=TXN; ORFNames=Ri
Homo sapiens (Human)
Eukaryota; Metazoa; (
Mammalia; Eutheria; l
                                                                                                                                                                                                                                                                                                           Q8W710 9CAUD
Q8W710;
01-MAR-2002 (
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Q5T936;
01-FEB-2005 (
01-FEB-2005 (
01-FEB-2005 (
                NUCLEOTIDE SEQUENCE.

MEDILINB=21972319; PubMed=11976141;

DOI=10.1128/AEM.68.5.2589-2594.2002;

Chen F., Lu J.;

"Genomic sequence and evolution of multiple of the control of the contro
                                                                                                                                                                        Cyanophage P60.
Viruses; dsDNA viruses,
NCBI_TaxID=151528;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GO; GÖ:0005489; F:electron transporter activity; GO; GO:0006118; P:electron transport; IEA. InterPro; IPR006662; Thiored. InterPro; IPR006663; Thioredox dom2. InterPro; IPR006663; Thioredox IPR006663; ThioredoxIn-like. Pfam; PF00085; ThioredoxIn-like. Pfam; PF00085; ThioredoxIn. PRINTS; PR00421; THIOREDOXIN. PROSITE; PS00194; THIOREDOXIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (MAY-2005)
EMBL; AL158158; CAII:
SMR; Q5T936; 1-85.
                                                                                                                                                                                                                                               Thioredoxin
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Intermediate filament; Keratin.
NON TER 85
SEQÜENCE 85 AA; 9314 MW; 40
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Microbiol.
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lysogenic phages.";
biol. 68:2589-2594(2002)
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Catarrhini; Homi
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RESULT 59
THIO2 DICCI
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THIO2 DICDI
P29446;
01-APR-1993
01-APR-1993
10-MAY-2005
Name=trxB; Synonyms=trx2;
Dictyostelium discoideum (Slime mold).
Eukaryota; Mycetozoa; Dictyosteliida;
                                                                  Thioredoxin
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Query Match
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Q9C922;
01-JUN-2001 (
01-JUN-2001 (
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Town C.D., Kaul S.;
Town C.D., Kaul S.;
Submitted (JAN-2001) to the
EMBL; AC019018; AAG52290.1;
HSSP, P23400; 1DBY.
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                                                                                                                                                                                                                       Pfam; PF00085; Thioredoxin; 1. PRINTS; PR00421; THIOREDOXIN. PROSITE; PS00194; THIOREDOXIN;
                                                                                                                                                                                                                                                                   GO; GO:0005489; F:electron transporter activity; GO; GO:0006118; P:electron transport; IEA.

InterPro; IPR006662; Thiored.

InterPro; IPR006663; Thioredox dom2.

InterPro; IPR012336; ThioredoxIn-like.

InterPro; IPR012335; ThioredoxIn fold.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
Lin X., Kaul S., Town C.D
Maiti R., Ronning C.M., K
Maiti R., Bowman C.
Barnstead M.E., Bowman C.
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NCBI_TaxID=3702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;
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01-JUN-2001 (TrEMBLrel. 17, Last sequence up
01-MAR-2004 (TrEMBLrel. 26, Last annotation
Putative thioredoxin; 109829-109566.
                                                                                                                                                                                                           SEQUENCE
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STANDARD;
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g C.M., Koo H., Fujii C.Y., Utterback T.R.,
Bowman C.L., White O., Nierman W.C., Fraser
.99) to the EMBL/GenBank/DDBJ databases.
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100.0%; Pr
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Pred. No. 4.4
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Pred. No. 4.4
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88
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(Rel. 25, Created)
(Rel. 25, Last sequence up
(Rel. 47, Last annotation
2 (Fragment).

update)

update)

Dictyostelium.

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Wetterauer B., Jacquot J.-P., Veron M.;
"Thioredoxins from Dictyostelium discoideum are a developmentally regulated multigene family.";
J. Biol. Chem. 267:9895-9904(1992).
-i- FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulfiand catalyzes dithiol-disulfide exchange reactions.
-i- SIMILARITY: Belongs to the thioredoxin family.
-i- SIMILARITY: Contains 1 thioredoxin domain.
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ACT_SITE
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InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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PIR; B46264; B46264.
HSSP; P10599; 1ERV.
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NUCLEOTIDE SEQUENCE.
STRAIN-ATCC 43049;
PubMed=15520287; DOI=10.1101/gr.2700304;
Baliga N.S., Bonneau R., Facciotti M.T.,
Deutsch E.W., Shannon P., Chiu Y., Weng F
Date S.V., Marcotte E., Hood L., Ng W.V.;
                                                                                                                                                                                                                                                                                                                                                                                                                           01-FEB-2005 (TrEMBLrel. 29,
01-FEB-2005 (TrEMBLrel. 29,
01-FEB-2005 (TrEMBLrel. 29,
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Q5V0J2;
01-FEB-2005 (TrEMBLrel. 2
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                                                                                                                                                                                                                                                                     Archaea; Euryarchaeota; Halobacteria; Halobacteriales;
Halobacteriaceae; Haloarcula.
                                                                                                                                                                                                                                                                                                                               Name=trxA3; OrderedLocusNames=rrnAC2110;
Haloarcula marismortui (Halobacterium ma
                                                                                                                                                                                                                                                                                                                                                                                                      Thioredoxin.
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Nucleophile (By similarity). 
Nucleophile (By similarity). 
Deprotonates C-terminal active site Cys
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RESULT 61
QBPZQ6_METMA ID
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QBPZQ6;
DT 01-CCT-2002 (1)
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DE ThioredLocusNe
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Best Local S
Matches 4
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HSSP; P80579; INW2.
GO; GO:0005489; F:electron transporter a
GO; GO:0005489; F:electron transport; IE
InterPro; IPR006662; Thiored.
InterPro; IPR005663; Thioredoxin.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin, 1.
PRINTS; PR00421; THIOREDOXIN.
TIGREAM6; TIGR0168; thioredoxin; 1.
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InterPro; IPR005746; Thioredoxin.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR006663; Thioredox dom2.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
TIGREAMS; TIGR01068; Thioredoxin; 1.
Complete proteome.
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Martinez-Arias R., Henne A., Wiezer A., Baeumer S.
Brueggemann H., Lienard T., Christmann A., Boemeck
Bhattacharyya A., Lykidis A., Overbeek R., Klenk H
Fritz H.-J., Gottschalk G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Methanosarcina mazei (Methanosarcina fri
Archaea; Euryarchaeota; Methanomicrobia;
Methanosarcinaceae; Methanosarcina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genome Res. 14:2221-2234(2004).
EMBL; AY596297; AAV46961.1; -; Genomic
GO; GO:00061489; F:electron transporter
GO; GO:0006118; P:electron transport; 1
                                                                                                                                                                            PRINTS; PR00421; THIOREDOXIN.
TIGREPAMS; TIGR01068; thioredoxi
PROSITE; PS00194; THIOREDOXIN;
Complete proceome.
SEQUENCE 88 AA; 9914 MW; D4
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EMBL; AE013269; AAM30132.1; -;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=Goe1 / Go1 / ATCC BAA-199 / DSM 3647 / OCM MEDLINE=22120827; PubMed=12125824;
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                                                                                                          Similarity
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between Bacteria and Archaea.
Microbiol. Biotechnol. 4:453-461(2002)
013269; AAM30132.1; -; Genomic_DNA.
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Pred. No. 4.4
D; Mismatches
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Pred. No.
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                                                                                                                 4.4e+02;
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                                                                                                                                      DB 2;
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                                                                                                                                    Length 88;
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Gunsalus R.
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Best Local S
Matches 4
                                                                  STRAIN=ATCC 23344;

PubMed=15377793; DOI=10.1073/pnas.0403306101;

Nierman W.C., DeShazer D., Kim H.S., Tettelin H., Nelson K.E., Rierman W.C., DeShazer D., Kim H.S., Tettelin H., Nelson K.E., Feldblyum T.V., Ulrich R.L., Romning C.M., Brinkac L.M., Daugherty S.C., Davidsen T.D., DeBoy R.T., Dimitrov G., Dodson Durkin A.S., Gwinn M.L., Haft D.H., Khouri H.M., Kholnay J.F., Madupu R., Mohammoud Y., Nelson W.C., Radune D., Romero C.M., Madupu R., Wohammoud Y., Nelson W.C., Radune D., Romero C.M., Sarria S., Selengut J., Shamblin C., Sullivan S.A., White O., Sarria S., Selengut J., Shamblin C., Sullivan S.A., White O., Safar N., Zhou L., Fraser C.M.;

"Structural flexibility in the Burkholderia mallei genome.";

Proc. Natl. Acad. Sci. U.S.A. 101:14246-14251(2004).

EMBL, CP000010; AAUS0201.1; -; Genomic_DNA.
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NON_TER
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Submitted (FEB-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF047604; AAC04362.1; -; Genomic_DNA.
GO; GO:0005576; C:extracellular region; IEA.
GO; GO:000579; F:hormone activity; IEA.
GO; GO:0005213; F:structural constituent of chorion (sensu InterPro; IFR005208; Cys_knot.
InterPro; IFR001545; Gly_hormoneB.
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25-OCT-2004 (TrEMBLrel.
25-OCT-2004 (TrEMBLrel.
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Pfam; PF00007; Gys knot; 1.

SMART; SM00068; GHB; 1.

PROSITE; PS00689; GLYCO_HORMONE_BETA_2;
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Q71UV5;
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Equus caballus (Horse).
Equus caballus (Horse).
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Dut-2004 (TrEMBLrel. 27, Li
Luteinizing hormone/chorionic
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NCBI_TaxID=13373;
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Bacteria; Proteobacteria; Betaproteobacteria; Burkholderiales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hypothetical protein.
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Mammalia; Eutheria;
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BURMA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 4; Conserv
  proteome;
88 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Burkholderia.
  Hypothetical protein.
9445 MW; 74D511CDE7B2ECD6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9153 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Laurasiatheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88
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27, Last sequence update)
27, Last sequence update)
27, Last annotation update)
prionic gonadotrophin beta-subunit
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Last sequence update)
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Pred.
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heria; Perissodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 2;
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Equidae; Equus.
                                                                                                                                                                                                                                                        Dodson R.J.,
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RESULT OF LONG MAN AND RAA MAN
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                                                                                                    .618 EQUZE PRELIMINARY; PR
O46618; EQUZE PRELIMINARY; PR
O46618; O1-JUN-1998 (TrEMBLrel. 06, Creat)
O1-JUN-1998 (TrEMBLrel. 24, Last
f 01-JUN-2003 (TrEMBLrel. 24, Last
f 01-JUN-2003 (TremBLrel. 24, Last
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Best Local Similarity
Matches: 4; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
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Q9HMDO;
01-MAR-2001 (TrEMBLrel. 16
01-MAR-2004 (TrEMBLrel. 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=20504483; PubMed=11016950; DOI=10.1073/pnas.190337797; Mg W.V., Kennedy S.P., Mahairas G.G., Berquist B., Pan M., Shukla H.D., Lasky S.R., Baliga N.S., Thorsson V., Sbrogna J., Swartzell S., Weir D., Hall J., Dahl T.A., Welti R., Goo Y.A., Leithauser B., Keller K., Cruz R., Danson M.J., Hough D.W., Maddocks D.G., Jablonski P.E., Krebs M.P., Angevine C.M., Dale H., Isenbarger T.A., Peck R.F., Pohlschroder M., Spudich J.L., Jung K., Isenbarger T.A., Peck R.F., Pohlschroder M., Spudich J.L., Jung K., Blam M., Freitas T., Hou S., Daniels C.J., Dennis P.P., Omer A.D., Ebhardt H., Lowe T.M., Liang P., Riley M., Hood L., DasSarma S.; "Genome sequence of Halobacterium species NRC-1."; Proc. Natl. Acad. Sci. U.S.A. 97:12176-12181 (2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Halobacterium salinarium (Halobacterium halobium).
Archaea; Euryarchaeota; Halobacteria; Halobacteriales;
Halobacteriaceae; Halobacterium.
NCBI_TaxID=2242;
Equus zebra hartmannae.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Laurasiatheria; Perissodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIR; E84409; E84409.
HSSP; P10599; IERV.
GO; GO:0005489; F:electron transporter activity;
GO; GO:00066118; P:electron transport; IEA.
GO; GO:0006810; P:transport, IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].
STRAIN=NRC-1 / ATCC 700922 / JCM 11081;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thioredoxin.
                                                                              Name=LH/CG-beta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRINTS; PR00421; THIOREDOXIN.
TIGRFAMs; TIGR01068; thioredo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=trxA2; OrderedLocusNames=VNG2600G;
                                                                                                 (Fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR006662; Thiored.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR006663; Thioredox_dom2
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HALSA
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1 (TrEMBLrel. 16,
4 (TrEMBLrel. 26,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10019 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thioredoxin;
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ic gonadotrophin beta-
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Pred. No. 4.4
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BDF97B3496F70BFF CRC64;
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                                                                                                                                   update)
beta-subunit
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Euteleostomi;
Equidae; Equu
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RESULT
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RESULT 67
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Matches 4
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Matches 4
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GO; GO:0005179; F:hormone activity; IEA.
GO; GO:0005213; F:structural constituent of
InterPro; IPR006208; Cys_knot.
InterPro; IPR001545; Gly_hormoneB.
Pfam; PF00007; Cys_knot; 1.
SMART; SM00068; GHB; 1.
PROSITE; PS00689; GLYCO_HORMONE_BETA_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fischer S. Veits J. Mey
Submitted (FBB-1998) to t
EMBL; AF047602; AAC04360.
HSSP; P01233; 1HCN.
GO; GO:0005576; C:extrace
GO; GO:0005179; F:hormone
GO; GO:0005213; F:structu
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19_EQUHM
046619_E
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SEQUENCE
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NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                         InterPro; IPR006208; Cys knot.
InterPro; IPR001545; Gly hormoneB.
Pfam; PF000007; Cys knot; 1.
SMART; SM00068; GHB; 1.
PROSITE; PS00689; GLYCO_HORMONE_BE
                                                                                                                                                                                                                                                                                                                                                                                                                                                      GO; G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE.
Fischer S., Veits J., Meyer
Submitted (FEB-1998) to the
EMBL; AF047603; AAC04361.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-1998 (TYEMBLrel. 06, Created)
01-JUN-1998 (TYEMBLrel. 06, Last sequence update)
01-JUN-2003 (TYEMBLrel. 24, Last annotation update)
Luteinizing hormone/chorionic gonadotrophin beta-subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9794;
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Mammalia; Eutheria; Laurasiatheria; Perissodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=LH/CG-beta;
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GO:0005576; C:extracellular region; IEA.
GO:0005179; F:bormone activity; IEA.
GO:0005213; F:structural constituent of
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tted (FEB-1998) to the EMBL/GenBank/DDBJ
AF047602; AAC04360.1; -; Genomic_DNA.
P01233; 1HCN.
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89
7 AA;
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9 AA;
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                                                                                                                                                                                                                                                                                                                           GLYCO_HORMONE_BETA_2;
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89
9325 MW;
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89
9369 MW;
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EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>.</u>
                                                                                                                                                                Score 31; DB 2; 1
Pred. No. 4.4e+02;
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Best Local (
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13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
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Q4HCJ2;
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13-SEP-2005 ('
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Name=UF-hep 2;
Paralichthys olivaceus (Japanese flounder).

Paralichthys olivaceus (Japanese flounder).

Paralichthys olivaceus (Japanese flounder).

Paralichthys olivaceus (Japanese flounder).

Actinopterygii; Percomorpha; Puteleostei; Neoteleostei;

Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;

Pleuronectoidei; Paralichthyidae; Paralichthys.

NCBI TaxID=8255;
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Hypothetical protein.
SEQUENCE 89 AA; 97
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STRAIN-DSM 11300;
US DOE Joint Genome Institute (JGI-ORNL);
Larimer F., Land M.;
"Annotation of the draft genome assembly of the Management of the draft genome assembly of the EMBL/GenBank/)
Submitted (JUN-2005) to the EMBL/GenBank/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Copeland A., Lucas S., Lapidus A., Barry K.,
Hammon N., Israni S., Pitluck S., Richardson
"Sequencing of the draft genome assembly of I
DSM 11300.";
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                                                                                                                  olivaceus.";
Submilted (JAN-2005) to the EMBL/GenBank/DDBJ
EMBL; AB199063; BAE06235.1; -; Genomic DNA.
SEQUENCE 89 AA; 9729 MW; 04CBB16152953F00
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STRAIN=DSM 11300;
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L; AAHE01000001; F
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                                                                                                            tted \.; AB198063; BAE
89 AA;
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                                       Conservative
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RESULT 70
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ID Q7R5Z2,
AC Q7R5Z2,
AC Q7R5Z2,
DT 01-MAR-2004 (TYEMBLYE1. 26,
DE GLP 81 67830 68102.
OS GIATGIA lamblia ATCC 50803.
OC BUKARYOCE; Diplomonadida; H
OX NCBI_TAXID=184922;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RC STRAIN=WB C6;
RA MOYRISON H.G., MCARThur A.G
RA Olsen G.J., Sogin M.L.;
"DYAST sequence of the Glarv
RL Submitted (MAR-2003) to the
CC EMBL/GenBank/DDJ whole
CC EMBL/GenBank/DDJ whole
CC EMBL/AACBO100003; EAAA270;
SQ SEQUENCE 90 AA; 10714 MW
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Search completed: February 23, 2006, 00:41:45 Job time: 177.5 secs
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Best Local Similarity 100
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                                                     Olsen G.J., Sogin M.L.;
"Draft sequence of the Giardia lamblia genome.";
submitted (MAR-2003) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DBJ whole genome shotgun (WGS) entry which is
preliminary data.
EMBL; AACB0100003; EAA42709.1; -; Genomic DNA.
EMBL; AACB0100003; EAA42709.1; -; Genomic DNA.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Actinopterygil; Neopterygil; Teleostei; Buteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygil; Percomorpha; Pleuronectiformes;
Pleuronectoidei; Paralichthyidae; Paralichthys.

NCBI TaxID=8255;
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Giardia lambTia ATCC 50803.

Bukaryota; Diplomonadida; Hexamitidae; Giardiinae; Giardia.

MCBI TaxID=184922;
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Submitted (JAN-2005) to the EMBL/GenBank/DDBJ databases.
EMBL; AB198061; BAE06233.1; -; mRNA.
SEQUENCE 89 AA; 9771 MW; D9AE7173C29CB412 CRC64;
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01-MAR-2004 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
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Hirono I., Nozaki R., Ono Y., Kurobe T., Ohira T., Aoki T.;
"Two different types of hepcidins from Japanese flounder Paralichthys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hepcidin 2.
Name=JF-hep 2;
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Pred. No. 4.4e+02;
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Pred. No. 4.5e+0;
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RESULT 1
US-07-820-688A-5
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Patent No.
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/820,688A
FILING DATE: 1920324
CLASSIFICATION DATA:
APPLICATION UMBER: PCT/NL91/00101
FILING DATE: 18-JUN-1991
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: PCT/NL91/00101
FILING DATE: 18-JUN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 90201598.1
FILING DATE: 18-JUN-1990
ATTORNEY/AGENT INFORMATION:
NAME: RAE-VENTER PH.D., BARBARA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: NAN, HELE
APPLICANT: KATTEVILD
APPLICANT: PALISSER,
TITLE OF INVENTION:
TITLE OF INVENTION:
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CITY: PALO ALTO
STATE: CALIFORNIA
COUNTRY: USA
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AV-GAY, YOSSEF
NAN, HELENA M.
KATTEVILDER, ALFRED
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VENTION: AN OXIDO REDUCTASE ENZYME SYSTEM
VENTION: OBTAINABLE FROM P. CHRYSOGENUM, THE SET OF GENES ENCODING
VENTION: THE SAME AND THE USE OF OXIDO REDUCTASE ENZYME SYSTEMS OR
VENTION: GENES ENCODING THE SAME FOR INCREASING ANTIBIOTIC PRODUCTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BOVENBERG, ROELOF A.L. SCHREIBER, RACHEL
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VAN DER VOORT, LUCIA H
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US-09-562-702A-30
US-09-561-818A-26
US-10-037-182-18
US-09-949-016-7247
US-09-949-016-53B-3
US-08-477-451-12
US-09-561-709B-1
US-09-561-818A-12
US-09-561-818A-12
US-09-949-016-9938
US-08-479-722B-2
US-09-949-016-9938
US-08-479-722B-2
US-09-951-818A-12
US-09-951-818A-12
US-09-951-818A-12
US-09-951-818A-12
US-09-951-818A-12
US-09-951-818A-12
US-09-951-685-6
PCT-US95-08251-18
US-08-347-594A-2
US-08-347-594A-2
US-08-347-594A-2
US-08-347-594A-2
US-08-347-594A-2
US-08-347-594A-2
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US-09-562-702A-32
US-09-561-818A-28
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US-10-188-495-67
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ALIGNMENTS
                                                                                                                                                                                                         #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Appli
Sequence 67, Appl
Sequence 67, Appl
Sequence 28, Appl
Sequence 28, Appl
Sequence 30, Appl
Sequence 30, Appl
Sequence 18, Appl
Sequence 18, Appli
Sequence 12, Appli
Sequence 1, Appli
Sequence 10, Appli
Sequence 10, Appli
Sequence 5938, Appli
Sequence 2, Appli
Sequence 3, Appli
Sequence 3, Appli
Sequence 6, Appli
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US-08-274-043B-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 32750
REFERENCE/DOCKET NUMBER: GRO
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-7622
TELEPAX: (415) 857-663
TELEX: 380816 COOLEY PA
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Loc
Matches
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APPLICANT: Aharono
APPLICANT: Van Der
APPLICANT: Cohen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
                                                  FILING DATE: 12-JULY-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/820,688
FILING DATE: 24-MARCH-1992
APPLICATION NUMBER: PCT/ML91/00101
FILING DATE: 18-JUNE-1991
APPLICATION NUMBER: EPO 90201598.1
FILING DATE: 18-JUNE-1990
ATTORNEY/AGENT INFORMATION:
NUMBER/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Kattevilde
APPLICANT: Pallissa,
TITLE OF INVENTION: F
TITLE OF INVENTION: F
TITLE OF INVENTION: C
TITLE OF INVENTION: C
TITLE OF INVENTION: C
                                                                                                                                                                                                         ZIP: 94306
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" diskette
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/274,043B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: protein FEATURE:
                                                                                                                                                                                                                                                                                                                                                                 STREET: P.O. Box
CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 4 amino a TYPE: AMINO ACID TOPOLOGY: linear
                         NAME: Rae-Venter, Barbara REGISTRATION NUMBER: 32,750
                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5, Application US/08274043B
o. 5652132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Argaman, Anat
Av-Gay, Yossesf
Nan, Helena M.
Kattevilder, Alfred
Pallissa, Harriet
Pallissa, Tarvido rec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 amino acids
                                                                                                                                                                                                                                                                                                                                                   U.S.A.
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Schreiber, Rachel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Van Der Voort, Lucia H.
Cohen, Gerald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Aharonowitz, Yair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           redox sequence typical for thioredoxins 1...4
                                                                                                                                                                                                                                                                                                                                                                                                   Law Offices of Barbara Rae-Venter O. Box 60039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 31;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               An oxido reductase enzyme system obtainable from P. chrysogenum, the set of genes encoding the same and the oxido reductase enzyme systems or genes encoding the same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   increasing antibiotic production
                                                                                         EPO 90201598.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roelof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,</u>
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           GBRO-024/01US
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TELECOMMUNICATION INFORMATION:

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Best Local Similarity
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/274,043
FILING DATE: 12-JULY-1994
APPLICATION NUMBER: US 07/820,688
FILING DATE: 24-MARCH-1992
APPLICATION NUMBER: PCT/NL91/000101
FILING DATE: 18-JUNE-1991
APPLICATION NUMBER: EP 90201598.1
APPLICATION NUMBER: EP 90201598.1
FILING DATE: 18-JUNE-1990
ATTORNEY/AGENT INFORMATION:
NAME: Rae-Venter, Barbara
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: GBRO-024/02US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (415) 926-6200
TELEPAX: (415) 854-3713
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
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MEDIUM TYPE: 3.5" diskette
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/386,729A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Palliss:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                          FILING DATE: 10-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION:
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TOPOLOGY: linear
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2882 Sand Hill Road,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pallissa, Harriet
VENTION: An oxido reductase enzyme system
vention: obtainable from P. chrysogenum, from the use of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Argaman, Anat
Av-Gay, Yossef
Nan, Helena M.
Kattevilder, Alfred
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Cohen, Gerald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bovenberg, Roelof A. L. Schreiber, Rachel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             redox sequence typical for thioredoxins 1..4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gotshal & Manges
4 Hill Road, Ste.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    obtainable from P. chrysogenum, the set of genes encoding the same and the use of oxido reductase enzyme systems or genes encoding the same for increasing antibiotic production
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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     GBRO-024/02US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 31; DB 1;
Pred. No. 4.6e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                       Query Match
Best Local Similarity
Matches 4; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3, Application US/08687276
Patent No. 5910435
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                             COMPUTER: IBM PC COMPATIBLE
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Vei
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/687,276
FILING DATE: 25-JUL-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27,386
REGISTRATION NUMBER: 960296.939:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (415) 926-62
TELEFAX: (415) 854-3713
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                    TELEFAX: 608-251-9166
INFORMATION FOR SEQ ID NO:
                                                                                                                               TOPOLOGY: lir
MOLECULE TYPE:
FRAGMENT TYPE:
                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Quarles &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Raines, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 53703
                                                                                                                                                                           STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                   TELEPHONE:
                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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                            2 CGPC 5
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                                                                                                                                                                                           amino acids
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GY: linear
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                                                       Conservative
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                                                                                                                                                             linear
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                                                                                                                                                                                                                                                                     608-251-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Quarles & Brady
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (415) 926-6200
                                                                                                                               peptide
internal
                                                                      100.0%;
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                                                                     Score 31; DB 1;
Pred. No. 4.6e+05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 31; DB 1;
Pred. No. 4.6e+05;
Nismatches 0;
                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 4;
                                                                                   Length 4;
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                                                       Gaps
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/883,804
FILING DATE:
CLASSIFICATION NUMBER: US/08/883,804
FILING DATE:
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Connell, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2879-40
TELECOMMUNICATION: INFORMATION:
TELEPHONE: 303/863-9700
TELECPHONE: 303/863-9700
TELECPAX: 303/863-9700
TELECPAX: 303/863-023
INFORMATION FOR SEQ ID NO: 2:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-504-538A-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21, Application US/08504538A Patent No. 6004746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 31; DB 1; Length 4; Best Local Similarity 100.0%; Pred. No. 4.6e+05; Matches 4; Conservative 0; Mismatches 0; Indels
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APPLICANT: White, Carl W.
APPLICANT: Das, Kumuda C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
ZIP: 02110-2214
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                             APPLICANT: Brent, Roger
APPLICANT: McCoy, John M.
APPLICANT: McCoy, John M.
APPLICANT: Jessen, Timm H.
TITLE OF INVENTION: INTERACTION TRAP SYSTEMS FOR DETECTING
TITLE OF INVENTION: PROTEIN INTERACTIONS
                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: Clark & Elbing LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheridan Ross P.C.
STREET: 1700 Lincoln Street, Suite 3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: USE OF THIOREDOXIN-LIKE MOLECULES FOR TITLE OF INVENTION: INDUCTION OF MASOD TO TREAT OXIDATIVE DAMAGE NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
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COUNTRY: U.
ZIP: 80203
                                                                                                                                                    COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acids
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                                                                                                                                                                                                                            176 Federal Street
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                                              GENERAL INFORMATION:
APPLICANT: BESCKWITH, JONATHAN
APPLICANT: ASLUND, FREDRIK
APPLICANT: BESSETTE, PAUL H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin version 3.0
SEQ ID NO 1
LENGTH: 4
TYPE: PRT
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GENERAL INFORMATION:
                                                                                                                     Sequence 2, Application US/09679705 Patent No. 6872563
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Best Local &
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FILING DATE: 07/20/95
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/278,082
FILING DATE: 07/20/94
ATTORNEY/AGENT INFORMATION:
NAME: Paul T. Clark
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 428-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: RABENSTEIN, DALLAS
APPLICANT: SHI, TIESHENG
TITLE OF INVENTION: CHEMICAL REAGENTS FOR FORMATION OF DISULFIDE BONDS IN PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 407T-891100US
CURRENT APPLICATION NUMBER: US/09/579,883A
CURRENT FILING DATE: 2000-05-26
                  APPLICANT:
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 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: synthetic peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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Local Similarity 100.0%; Pred. No. 4.6e+09
hes 4; Conservative 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
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RITZ, DANIEL
LIM, JACKIE EUN-AH
                                 GEORGIOU,
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100.0%; Pred. No. 4.6e+05;
ative 0; Mismatches 0;
                                 GEORGE
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RESULT 9
US-08-883-804-4
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; OTHER INFORMATION: Description of Artificial Sequence: Synthetic;
; OTHER INFORMATION: variant motif
US-09-679-705-2
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Best Local Similarity
  Query Match
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APPLICANT: White, Carl W.
APPLICANT: Das, Kumuda C.
TITLE OF INVENTION: USE OF
TITLE OF INVENTION: INDUCTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/166,044
PRIOR FILING DATE: 1999-11-17
NUMBER OF SEQ ID NOS: 24
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PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/163,939
PRIOR FILING DATE: 1999-11-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/679,705
CURRENT FILING DATE: 2000-10-05
                                                                                                                                                                                                                 CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: CONNELL, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2879
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR PRODUCTION OF DISULFIDE TITLE OF INVENTION: BOND CONTAINING PROTEINS IN HOST CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
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                                                                                                                                                           INFORMATION FOR SEQ ID NO:
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tent No. 5985261
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                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/883,804
                                                              MOLECULE TYPE:
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                                                                                  TOPOLOGY:
                                                                                                                     LENGTH:
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                                                                                amino acid
GY: linear
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1700 Lincoln Street, Suite 3500
                                                                                                                     6 amino acids
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  100.0%;
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  Score 31;
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DB 1;
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RESULT 11
US-08-627-907A-6
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Best Local Similarity
Watches 4; Conserve
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US-08-883-804-5
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Patent No. 6060302
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Patent No.
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                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/883,804
FILING DATE:
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO:
                                                                       APPLICANT: HIRANO, NAOTO
APPLICANT: HIRAI, Hisamaru
TITLE OF INVENTION: HUMAN PHOSPHOLIPASE C-ALPHA AND DNA
TITLE OF INVENTION: SEQUENCE ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: CONNELL, GATY J.
REGISTRATION UNMBER: 32,020
REFERENCE/DOCKET NUMBER: 28
REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                             NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheridan Ross P.C.
STREET: 1700 Lincoln Street, Suite 3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: White, Carl W. APPLICANT: Das, Kumuda C. TITLE OF INVENTION: USE O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
TOPOLOGY:
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Pred. No. 4.6e+05;
D; Mismatches 0;
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                                                                                                ; OTHER INFORMATION: S. US-07-901-713A-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 22-SEP-1994
ATTORNEY/AGENT INFORMATION:
NAME: HALEY Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: SHON-10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
TELEFAX: (212) 596-9000
TELEFAX: (212) 596-9000
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                               Sequence 3, Application US/07901713A

Patent No. 6291205

GENERAL INFORMATION:

APPLICANT: Tuite, Michael F.
APPLICANT: Freedman, Robert B.
APPLICANT: Schultz, Loren D.
APPLICANT: Schultz, Loren D.
APPLICANT: Montgomery, Donna L.
APPLICANT: Montgomery, Donna L.
APPLICANT: Bllis, Ronald W.
TITLE OF INVENTION: METHOD FOR INCREASING PRODUCTION OF
TITLE OF INVENTION: CEREVISIAE

FILE REFERENCE: 18469
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Best Local S
                                                                                                                                                                                                                                     SEQ ID NO 3
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                                                  Query Match
                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/07/901,713A
CURRENT FILING DATE: 1992-06-12
NUMBER OF SEO ID NOS: 23
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA
APPLICATION NUMBER: US/08/627,907A
                                                                                                                                                                                                                 ENGTH: 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: JP 5-238402
FILING DATE: 24-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/JP94/01572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          y Match 100.0%; Score 31; DB 2; 1
Local Similarity 100.0%; Pred. No. 4.6e+05;
hes 4; Conservative 0; Mismatches 0;
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ZIP: 10020
y Match 100.0%; Score 31; DB 2; Length 6;
Local Similarity 100.0%; Pred. No. 4.6e+05;
hes 4; Conservative 0; Mismatches 0; Indels
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                                                                                                                      cerevisiae PDI thioredoxin active site
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       Gaps
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; LENGTH: 8 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-571-985-27
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US-08-571-985-27
Sequence 27, Application US/08571985
Patent No. 5783557
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US-09-116-766-27
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Matches
                                                                                                                                                                         Sequence 27, Applicat Patent No. 5968898 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/571,985
FILING DATE:
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Burstein, Yigal
APPLICANT: Trainin, Nathan
APPLICANT: Rycus, Avigail
TITLE OF INVENTION: THF-gamma2 Analogs and Pharmaceutical
TITLE OF INVENTION: Compositions Comprising Them
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                         APPLICANT: Burstein, Yigal
APPLICANT: Trainin, Nathan
APPLICANT: Rycus, Avigail
TITLE OF INVENTION: THF-gamma2 Analogs and Pharmaceutical
TITLE OF INVENTION: Compositions Comprising Them
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 01-JUL-1993
ATTORNEY/AGENT INFORMATION:
NAME: Kohn, Kenneth I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 30,955
REFERENCE/DOCKET NUMBER: 17
TELECOMMUNICATION INFORMATION:
TELEPHONE: (810) 539-5050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: IL 106214
                                   NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 30500 No. 5783:
CITY: Farmington Hills
                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP:
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   STREET:
                     DDRESSEE:
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Y: USA
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30500 No. 5783557thwestern Hwy., Suite 410
E: Kohn & Associates
30500 No. 5968898thwestern Hwy., Suite 410
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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100.0%; Pred. No. 4.6e+05;
tive 0; Mismatches 0;
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COUNTRY:

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (248) 539-5050
TELEPAX: (248) 539-5055
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
APPLICANT:
APPLICATION NUMBER: US/08/836,480
FILING DATE: 14-APR-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: SE 95\403256-8
FILING DATE: 14-OCT-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                          COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                 ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Karabelas, Kostas
APPLICANT: Lindvall, Magnus
APPLICANT: Sarnstrand, Bengt
TITLE OF INVENTION: NEW PEPTIDES WITH IMMUNOMODULATORY EFFECTS
NUMBER OF SEQUENCES: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 11 106214
FILING DATE: 01-JUL-1993
ATTORNEY/AGENT INFORMATION:
NAME: MODISOMERY, 11ene N.
REGISTRATION NUMBER: 38,972
REFERENCE/DOCKET NUMBER: 2163.00050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TYPE: linear
                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                              CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
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Erickson, Tomas
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Pred. No. 4.6e+05;
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; MOLECULE TYPE: peptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 34:
US-09-562-897-34
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US-09-562-897-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: peptide US-08-836-480-34
Query Match 100.0%; Score 31; DB 2; Best Local Similarity 100.0%; Pred. No. 4.6e+05; Matches 4; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 34, Application Patent No. 6228374
GENERAL INFORMATION:
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Best Local (
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                                                                                                                                                                                                                 TELEFAX: 617-542-8906 INFORMATION FOR SEQ ID NO: 34: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 06:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5970
TELEPHAX: 617-542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNMBER: US/09/562,897
FILING DATE: 01-May-2000
CLASSIFICATION: CUNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/836,480
FILING DATE: <UNKnown>
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION UMBER: 34,819
REFERENCE/DOCKET NUMBER: 06275/036001
TELECOMMUNICATION INFORMATION:
TETILEDYNAM: 617-642-6070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    y Match 100.0%; Score 31; DB 2; Local Similarity 100.0%; Pred. No. 4.6e+05; Les 4; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 8 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sarnstrand, Bengt
TITLE OF INVENTION: NEW PEPTIDES WITH IMMUNOMODULATORY EFFECTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bergstrand, Hakan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 CGPC 5
                                                                                                                                                    TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                    TELEPHONE: 617-542-5070
                                                                                                                                                                                                 LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09562897
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Karabelas,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lindvall, Magnus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Erickson, Tomas
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                                           Length 8;
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US-08-750-142B-54
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US-09-160-513-217
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                                                                                                                                           Sequence 217, Application US/09160513 Patent No. 6410775
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Best Local Similarity
Matches 4; Conser
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/SE96/00:
FILING DATE: 22-MAR-1996
APPLICATION NUMBER: SE9501067-4
FILING DATE: 24-MAR-1995
ATTORNEY/AGENT INFORWATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FASTSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/750,142B
FILING DATE: 25-NOV-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Bergstrand, Hakan
APPLICANT: Eriksson, Tomas
APPLICANT: Lindvall, Magnus
APPLICANT: Sarnstrand, Bengt
TITLE OF INVENTION: NEW PEPTIDES WITH
TITLE OF INVENTION: IMMUNOMODULATORY EFFECTS
             APPLICANT: Victoria, Edward J.
APPLICANT: Marquis, David M.
APPLICANT: Jones, David S.
APPLICANT: Yu, Lin
TITLE OF INVENTION: aPL IMMUNOREACTIVE PEPTIDES, CONJUGATES THEREOF AND METHODS NUMBER OF SEQUENCES: 225
                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 617-542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 06275/062001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
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                                                                                                                                                                                                                                                                                              2 CGPC 5
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Pred. No. 73;
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; ORGANISM: Rhizobium leguminosarum
US-09-230-041-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE: peptide US-09-160-513-217
                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 31
LENGTH: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 6465179
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 31, Application US/09230041A Patent No. 6465179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                Matches
                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                         APPLICANT: THIREOS, GEORGE
APPLICANT: KAPETZOPOULOS, DIMITRIS
APPLICANT: KAPETZOPOULOS, DIMITRIS
TITLE OF INVENTION: DAN ENCODING AN ARTHROPOD CHITIN SYNTH
CIURRENT EFERENCE: DNA ENCODING AN ARTHROPOD CHITIN SYNTH
CURRENT APPLICATION NUMBER: US/09/230,041A
CURRENT FILING DATE: 1999-01-19
EARLIER APPLICATION NUMBER: EP97108240.9
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 42
COUNTRIES OF SEQ ID NOS: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: CATHERINE M. POLIZZI
REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 25231-20061.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEPAX: (650) 494-0792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/160,513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 16 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 Match 100.0%; Score 31; DB Local Similarity 100.0%; Pred. No. 89; Mismatches 4; Conservative 0; Mismatches
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CLASSIFICATION:
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                                                                              4; Conservative
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                                                                              100.0%; Score 31; DB 2;
100.0%; Pred. No. 1.3e+02;
tive 0; Mismatches 0;
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RESULT 20 US-09-881-572A-25

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; Sequence 26, Application US/10044359
Patent No. 6740743
; GENERAL INFORMATION:
APPLICANT: Herrman, Rafael
APPLICANT: Wong, James F.
; APPLICANT: Lee, Jian-Ming
TITLE OF INVENTION: SCORPION TOXINS
FILE REFERENCE: BB1367 US NA
; CURRENT FILING DATE: 2002-01-11
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                                                                                                                                                                                                                                         US-10-044-359-26
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Best Local S
Matches 4
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PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn version 3.0
SEQ ID NO 25
LENGTH: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.0 SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/09881572A Patent No. 6762031
                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
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Patent No. 6762031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Bupp, Keith;
APPLICANT: University of Medicine and Dentistry of New Jersey
TITLE OF INVENTION: Targeting Viral Vectors to Specific Cells
FILE REFERENCE: 601-1-095
CURRENT APPLICATION NUMBER: US/09/881,572A
CURRENT FILING DATE: 2002-03-05
PRIOR APPLICATION NUMBER: 60/212,239
PRIOR APPLICATION STATE CONTROL OF SERVING DATE: 2000-06-16
NUMBER OF SERVING DATE: 2000-06-16
NUMBER OF SERVINGS: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Bupp, Keith,
APPLICANT: University of Medicine and Dentistry of New Jersey
TITLE OF INVENTION: Targeting Viral Vectors to Specific Cells
FILE REFERENCE: 601-1-095
CURRENT APPLICATION NUMBER: US/09/881,572A
CURRENT FILING DATE: 2002-03-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Roth, Monica; APPLICANT: Bupp, Keith;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Roth, Monica;
PRIOR APPLICATION NUMBER: 09/599,416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 30
TYPE: PRT
ORGANISM: amphotropic murine leukemia virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                    11
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                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 31; DB 2;
100.0%; Pred. No. 1.4e+02;
live 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 23
US-08-239-256-16
                                                                                                                        US-08-239-256-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Microsoft Office 97
SEQ ID NO 26
LENGTH: 35
                                                          Matches
                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%;
Best Local Similarity 100.0%;
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PRIOR APPLICATION NUMBER: 60/140,227
PRIOR:FILING DATE: 1999-06-22
NUMBER OF SEQ ID NOS: 28
                                                                                                                                                                                                  TELEX: 90-4030
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                              NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 2950
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
TELEFAX: (202) 887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: BOIME, IRVING
APPLICANT: MATZUK, MARTIN M.
APPLICANT: KEENE, JEFFREY L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: CTP EXTENDED FORM OF NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 2000 remay....
CITY: Washington, D.C.
                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/239,256 FILING DATE: 06-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                   TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                          IOPOLOGY:
                                                                                                                                                                                      LENGTH:
23
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CGPC 26
                                                                                                                                                                                     38 amino acids
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2000 Pennsylvania Ave. N.W.
                                                          Conservative
                                                                                                                                    linear
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                                                                         Score 31; DB 1;
Pred. No. 1.6e+02;
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Pred. No. 1.6e+02;
                                                            Mismatches
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                                                                                         Length 38;
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                                                            Gaps
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RESULT 24 US-09-244-583-20 ; Sequence 20, Application US/09244583

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US-09-244-583-20
                                                                                                                     US-09-579-420B-1
                                                                                                                                      RESULT 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/583,638
CURRENT FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: PCT/US98/26127
PRIOR FILING DATE: 1998-12-09
PRIOR APPLICATION NUMBER: 60/069,155
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR PPLICATION NUMBER: 60/069,687
PRIOR PILING DATE: 1997-12-12
PRIOR PILING DATE: 1997-12-12
PRIOR PILING DATE: 1997-12-12
PRIOR PILING DATE: 1997-12-12
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CURRENT FILING DATE: 1999-02-04
EARLIER APPLICATION UNMEER: 60/073,979
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 29
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 20
                Sequence 1, Application US/09579420B
Patent No. 6777534
GENERAL INFORMATION:
APPLICANT: Klagabrun, Michael
APPLICANT: Soker, Shay
                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 6635421
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.0%;
Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: BAIRD, ANDREW
APPLICANT: ANDREASON, GRAI
TITLE OF INVENTION: VOYEL FORMS OF THE ANGIOGENIC FACTOR
TITLE OF INVENTION: VASCULAR ENDOTHELIAL CELL GROWTH FACTILE REFERENCE: 240/086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: NEUROPILINS AND USE THEREOF IN METHODS FOR TITLE OF INVENTION: DIAGNOSIS AND PROGNOSIS OF CANCER FILE REFERENCE: 701039-48800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: KLAGSBRUN, MICHAEL
APPLICANT: SOKER, SHAY
TITLE OF INVENTION: PEPTIDE ANTAGONISTS OF VASCULAR ENDOTHELIAL GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 4:
TYPE: PRT
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MIAO, HUA-QUAN
TAKASHIMA, SEIJI
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Pred. No. 1.8e+02;
Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                Score 31; DB 2;
Pred. No. 1.8e+02;
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CURRENT APPLICATION NUMBER: US/09/579,420B
CURRENT FILING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: PCT/US98/26103
PRIOR FILING DATE: 1998-12-09
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR FILING DATE: 1997-12-12
PRIOR APPLICATION NUMBER: 60/069,155
PRIOR APPLICATION NUMBER: 60/069,155
PRIOR APPLICATION NUMBER: 50/069,155
PRIOR FILING DATE: 1997-12-09
NUMBER OF SEQ ID NOS: 19
SOFTWARE: PATENTIN VETBION 3.1
SEQ ID NO 1
                                                                                                                                                                                   RESULT 28
US-09-513-999C-7252
US-09-513-999C-7252, Application US/09513999C
; Patent No. 6783961
; Patent No. 6783961
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-579-420B-19
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US-09-579-420B-19
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Klagsbrun, Michael
APPLICANT: Soker, Shay
                                                                                                                                                                                 GENERAL INFORMATION:
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Best Local Similarity
                                           TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins Patent No. 6783961
FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local Similarity
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                                                                                                            APPLICANT: Dumas Milne Edwards, J.B
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
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CURRENT APPLICATION NUMBER: U90/579,420B
CURRENT FILING DATE: 2000-05-25
PRIOR APPLICATION NUMBER: PCT/US98/26103
PRIOR FILING DATE: 1998-12-09
PRIOR FILING DATE: 1998-12-12
PRIOR FILING DATE: 1997-12-12
PRIOR FILING DATE: 1997-12-09
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR FILING DATE: 1997-12-12
PRIOR FILING DATE: 1997-12-09
CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24
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ORGANISM: Homo sapiens
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Pred. No.
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; FEATURE:
, NAME/KEY: UNSURE
; LOCATION: 27
; OTHER INFORMATION: Xaa=Arg or Ser or Thr
US-09-513-999C-5904
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US-09-513-999C-5905
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Best Local S
Matches 4
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PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 5904
FEVERIFIED NO 5904
                                                                                                            Sequence 5905, Application US/09513999C Patent No. 6783961
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Glordano, J.Y.
APPLICANT: Glordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
FILE REFERENCE: 59 US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
CURRENT FILING DATE: 2000-02-24
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SOFTWARE: Patent.pm
SEQ ID NO 7252
                                                     TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins Patent No. 6783961
FILE REFERENCE: 59.US2.REG
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Best Local Similarity
CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24 PRIOR APPLICATION NUMBER: US 60/122,487
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TYPE: PRT
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Pred. No.
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2e+02;
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US-09-630-273-2
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Sequence 2, Application US/09630273

Patent No. 6521416

GENERAL INFORMATION:
Birken et al, Steven
TITLE OF INVENTION: Determination Of The Amount Of hith Beta Core Fragment
TITLE OF INVENTION: In A Sample From A Subject And Uses Thereof
FILE REFERENCE: 54204-A-PCT-US
FILE REFERENCE: 54204-A-PCT-US
CURRENT APPLICATION NUMBER: US/09/630,273
CURRENT FILING DATE: 2000-08-01
PRIOR APPLICATION NUMBER: PCT/US99/02279
PRIOR FILING DATE: 1999-02-03
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
APPLICANT: Wiegand, Roger C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15949)B
CURRENT APPLICATION NUMBER: US01-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR PILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 11828
LENGTH: 77
TYPE: NET
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NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 5905
LENGTH: 67
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Patent No. 6833447
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Best Local Similarity
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FEATURE:
NAME/KEY: UNSURE
LOCATION: 2
OTHER INFORMATION: Xaa=!
FEATURE:
NAME/KEY: UNSURE
LOCATION: 27
LOCATION: 27
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Local Similarity 100.0%; Pred. No. 2.7e+02;
hes 4; Conservative 0; Mismatches 0;
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Pred. No. 2.5e+02;
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RESULT 35
US-09-621-976-5629
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                                                                                                                                                                                                             ; TYPE: PRT ; ORGANISM: Chlamydia pneumoniae US-09-198-452A-697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 34
US-09-198-452A-697
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US-09-621-976-6168
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                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Griffais, R.
APPLICANT: Griffais, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999
FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 697
LENGTH: 102
TYPE: DET
                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 697, Application US/09198452A Patent No. 6559294
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Best Local Similarity
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SOFTWARE: Patent.pm
SEQ ID NO 6168
LENGTH: 89
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                                                                                                                                         Matches
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTs and Encoded Human Proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      63
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                                                                     CGPC 31
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; Pred. No. 2.8e+02;
red matches 0;
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APPLICANT: Jobert, S.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
FILE COFINUMION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Patent.pm
SEQ ID NO 5629
LENGTH: 102
TYPE: PRT
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GENERAL INFORMATION:
APPLICANT: Fang, Kathy S.
APPLICANT: Hanafusa, Hides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 5629, Application US/09621976 Patent No. 6639063
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                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION UNMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201 487-8800
TELEPAX: 201 343-1684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Dumas Milne Edwards, J.B.
                                                                                                                                      TELEX: 133521
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: SIGNAL
LOCATION: -35..-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 103 amino acids
                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/241,853 FILING DATE: 12-MAY-1994 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Hanafusa, Hidesaburo
TITLE OF INVENTION: A TRANSMEMBRANE TYROSINE PHOSPHATASE
TITLE OF INVENTION: AND METHODS OF USE THEREOF
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: New Jersey
                                      STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
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                  peptide
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100.0%; Pred. No. 3.3e+02;
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RESULT 38
US-09-583-110-3078
; Sequence 3078, Application US/09583110
; Patent No. 6699703
; GENERAL INFORMATION:
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                                                                                                                                                                Query Match
Best Local Similarity
Matches 4; Conserv
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GENERAL INFORMATION:
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/241,853
FILING DATE: 12-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-078
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201 487-5800
                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 103 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Hanafusa, Hidesaburo
TITLE OF INVENTION: A TRANSMEMBRANE TYROSINE PHOSPHATASE
TITLE OF INVENTION: AND METHODS OF USE THEREOF
NUMBER OF SEQUENCES: 37
                                                                                                                                                                                                                                                                HYPOTHETICAL:
                                                                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
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                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 201
TELEFAX: 133521
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FILING DATE: 02-MAY-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: UZIP: 07601
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STATE: New Jersey
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                                                                                                            52
                                                                                                                                                                                                                                                                                                            single
                                                                                                                                                                                100.0%; Score 31; DB 1; 100.0%; Pred. No. 3.3e+02;
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US-08-826-910-3
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APPLICANT: YODOI, JUNJI; UCHIDA, ATSUSHI; TAGAYA, YUTAKA AMITSUI, AKIRA, HIRAKAWA, TADASHI
FITLE OF INVENTION: METHOD FOR TRREATING CANCER THERAPY FRADIATION DAMAGE OR ARTERIOSCLEROSIS USING HUMAN ADF NUMBER OF SEQUENCES: 1

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5210073-1
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CURRENT APPLICATION NUMBER: US/09/583,110
CURRENT FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/107,433
PRIOR FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR APPLICATION NUMBER: US 60/051,553
PRIOR APPLICATION NUMBER: US 60/051,553
PRIOR FILING DATE: 1997-07-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS:
SEQ ID NO 3078
LENGTH: 104
TYPE: PRT
                                                                                                                                                                                                                                                                                                                              Sequence 3, Application US/08826910 Patent No. 5919657
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Best Local Similarity
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APPLICANT: Hillman, Jennifer L.
APPLICANT: Shah, Purvi
TITLE OF INVENTION: THIOREDOXIN-LIKE PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating
TITLE OF INVENTION: Pneumoniae for Diagnostics and Therapeutics
                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: FastSEQ for Wi
                                                                                              COUNTRY: UZIP: 94304
                                                                                                                                                    STREET: 3174 Por
CITY: Palo Alto
                                                                                                                                                                     ADDRESSEE: Incyte Pharmaceuticals, Inc
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                      STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 104
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    FastSEQ for Windows Version
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100.0%; Pred. No. 3.4e+02;
tive 0; Mismatches 0;
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US-08-826-910-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/08826910
Patent No. 5919657
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Shah, Purvi
TITLE OF INVENTION: THIOREDOXIN-LIKE PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                              ATTORNEY AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415.855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
IMMEDIATE SOURCE:
LIBRARY: GenBank
CLONE: 339649
                                                                                                                                                                                                         APPLICATION NUMBER: US/08/826,910
FILING DATE: Herewith
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                         OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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LENGTH: 105 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: Herewith
                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 31..
CITY: Palo Alto
CTATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 415-845-4166
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                                                  TELEFAX:
                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                 COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94304
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                                                    415-845-4166
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Pred. No. 3.4e+02;
); Mismatches 0;
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FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 8037
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LIBRARY: GenBa
CLONE: 453972
US-08-826-910-4
; LENGTH: 105
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-513-999C-8037
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US-09-601-144-68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 68
LENGTH: 105
TYPE: PRT
                                                                                                                                                                                                                        APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
                                                                                                                                                                                                                                                                                                                Sequence 8037, Application US/09513999C Patent No. 6783961 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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Patent No. 6566514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/073,196
PRIOR FILING DATE: 1998-01-30
NUMBER OF SEQ ID NOS: 74
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: OLIGONUCLEOTIDE SEQUENCES COMPLEMENTARY TO THIOREDOXIN TITLE OF INVENTION: AND THIOREDOXIN REDUCTASE GENES AND METHODS OF USING TITLE OF INVENTION: SAME TO MODULATE CELL GROWTH FILE REFERENCE: 683-112US-A CURRENT APPLICATION NUMBER: US/09/601,144

CURRENT FILING DATE: 2000-10-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Wright, Jim A. APPLICANT: Young, Aiping H. APPLICANT: Lee, Yoon S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: si
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100.0%; P.
ative 0;
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Pred. No.
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RESULT 44
US-08-386-729A-10
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                                                                ; MOLECULE TYPE: US-08-386-729A-10
                                                                                                                                             FILING DATE: 18-708-19-00
ATTORNEY/AGENT INFORMATION:
NAME: Rae-Venter, Barbara
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: GBRO-024/02US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 926-6200
TELEPAX: (415) 926-6200
TELEFAX: (415) 934-3713
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 106 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 4; Conserv
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PATENTINE PLOSS #1.00 Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/386,729A FILING DATE: 10-FEB-1995 CLASSIFICATION: 435
PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/274,043 FILING DATE: 12-UUY-1994 APPLICATION NUMBER: US 07/820,688 APPLICATION NUMBER: US 07/820,688 APPLICATION NUMBER: PCT/NL9/00101 FILING DATE: 24-WARCH-1992 APPLICATION NUMBER: PCT/NL9/00101 FILING DATE: 18-UUNE-1991 APPLICATION NUMBER: PCT/NL9/00101 FILING DATE: 18-UUNE-1991 APPLICATION NUMBER: EP 90201598.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Aharon
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MEDIUM TYPE: 3.5" ISSACTO
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: AV-Gay, Yossef
APPLICANT: Nan, Helena M.
APPLICANT: Nan, Helena M.
APPLICANT: Kattevilder, Alfred
APPLICANT: Value of National Control of Sense State
TITLE OF INVENTION: An oxido reductase enzyme system
TITLE OF INVENTION: obtainable from P. chrysogenum, the set of genes
TITLE OF INVENTION: encoding the same and the use of oxido reductase enzyme
TITLE OF INVENTION: systems or genes encoding the same for increasing antibiotic
TITLE OF INVENTION: production
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
Numbercer: Mail Cortehal & Manages
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APPLICANT:
APPLICANT:
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CITY: Menlo Park
STATE: CA
COUNTRY: U.S.A.
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                                                                                                        LENGTH: 106 amino acids
TYPE: amino acid
TOPOLOGY: linear
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Bovenberg, Roelof A. L. Schreiber, Rachel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Argaman, Anat
Av-Gay, Yossef
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100.0%;
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100.0%; Pred. No. 3.4e+02;
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Score 31; DB 1;
Pred. No. 3.4e+02;
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                    Length 106;
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APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Sleeman, Matthew
APPLICANT: Abernethy, Nevin
APPLICANT: Conrust, Rene
APPLICANT: Murison, Greg
ITILE OF INVENTION: Compounds isolated from stromal cells
ITILE OF INVENTION: Anand methods for their use
FILE REFERENCE: 11000.1037c1
CURRENT APPLICATION NUMBER: US/09/383,586
CURRENT FILING DATE: 1999-08-26
INUMBER OF SEQ ID NOS: 38
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 13
LENGTH: 106
TYPE: PRT
ODCANTION: Monica
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                                                                                                                               ; TYPE: PRT
; ORGANISM: Mouse
US-09-823-038A-13
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US-09-823-038A-13
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US-09-383-586-13
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US-09-383-586-13
                                                                                                                                                                                            APPLICANT: Kumble, Anand
APPLICANT: Murison, Greg
TITLE OF INVENTION: Compositions Isolated From Stromal Cells
TITLE OF INVENTION: and Methods For Their Use
FILE REFERENCE: 11000.1037c3
CURRENT APPLICATION NUMBER: US/09/823,038A
CURRENT FILING DATE: 2001-07-09
NUMBER OF SEQ ID NOS: 61
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
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                                                                Matches
                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 13, Application US/09823038A Patent No. 6797271
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Strachan,
                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                  LENGTH: 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            33 CGPC 36
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                                 2 CGPC 5
                                                              2h 100.0%; Score 31; DB 2; Similarity 100.0%; Pred. No. 3.4e+02. 4; Conservative 0; Mismatches 0.
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Abernethy, Nevin
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100.0%; Pred. No. 3.4e+02;
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                                                                                                 Length 106;
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                                                                Gaps
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RESULT 47
US-07-745-382-22
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US-07-921-848-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: CBerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI5188A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPHONE: (617) 876-5851
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                           Sequence 22, Applicati
Patent No. 5292646
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local (
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PAtentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/745,382
FILING DATE: 199.10814
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US 07/652,531
PILING DATE: 06-FEB-1991
ATTORNEY/AGENT INFORMATION:
NAME: CREAT LINEON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Like Molecules
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS: Genetics Institute, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                      APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Like Molecules
                                                                                                                                                                                    NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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CITY: Cambridge
                                                                                                                    STREET: 87 Cambridge
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                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                   ADDRESSEE:
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5292646
                                                                                                       Massachusetts
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                                                                                                                                            E: Genetics Institute, Inc. 87 CambridgePark Drive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 31; DB 1; ilarity 100.0%; Pred. No. 3.5e+02; Conservative 0; Mismatches 0;
                                                                                    U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-165-301A-22
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Best Local Similarity 100
Matches 4; Conservative
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APPLICATION NUMBER: US 07/652,531
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/745,382
APPLICATION NUMBER: US 07/745,382
PILING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORMATION:
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                                ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION UNMER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECHOME: (617) 876-1170
TELEPHONE: (617) 876-1170
TELEFAX: (617) 876-5851
TELEFAX: (617) 876-5851 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: b Cambridge CITY: Cambridge CTATE: Massachusetts
                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
FILING DATE:
CLASSIFICATION: 435
APPCDENT OF THE PROPERTY OF THE P
                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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LENGTH: 109 amino acid
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APPLICATION NUMBER: US,
FILING DATE: 19920728
CLASSIFICATION: 435
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TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
FORMATION FOR SEQ ID NO: 22:
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CORRESPONDENCE ADDRESS:
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SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A. ZIP: 02140
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REGISTRATION NUMBER: 31,822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 33 CGPC 36
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DiBlasio-Smith, Elizabeth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genetics Institute,
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US-08-165-301A-28

; Sequence 28, Application US/08165301A

; Patent No. 5646016
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                                                                                                                                                                                                                                                        JS-08-165-301A-26
                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: amino acid
                                                                                                                                                                                 Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC COMPATION
COMPUTER: PC COMPATION
COMPATION
COMPATION
COMPATION
COMPATION
COMPATION
CURRENT APPLICATION
CURRENT APPLICATION
APPLICATION
APPLICATION
APPLICATION
APPLICATION
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 87 Cambridges
CITY: Cambridge
STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                           33 CGPC 36
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                                                                                                                                 2 CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DiBlasio-Smith, Elizabeth
                                                                                                                                                                               100.0%; Score 31; I ilarity 100.0%; Pred. No. 3 Conservative 0; Mismatcher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genetics Institute, Inc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       GI
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                                                                                                                                                                                 Mismatches
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US-08-810-436-22
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                                                                                                                                                                                                                                                                                                                                                               Patent No. 6143524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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ZIP: UZING COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: YSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
                                                                                                                                                                                  APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
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ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: McCoy,
APPLICANT: DiBlasi
APPLICANT: Grant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
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REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
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                                                                                             STATE: M
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CITY: C
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STREET: 87 Cambr
CITY: Cambridge
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                                                                                               Massachusetts
': U.S.A.
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                                                                                                                                               E: Genetics Institute,
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Genetics Institute,
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 31; DB 1; illarity 100.0%; Pred. No. 3.5e+02; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                             DiBlasio-Smith,
                                                                                                                                                                                                                                                                                                                                  McCoy, John
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PEPTIDE AND PROTEIN FUSIONS TO THIOREDOXIN, THIOREDOXIN-LIKE HOLECULES
                                                                                                                                                                                                                                                                                                               Elizabeth
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SOFTWARE: Patentin Release #1.0, Version #1.25

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RESULT 53
US-08-810-436-26
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          ATTORNEY/AGENI A....

NAME: Meinert, M. C.

NAME: Meinert, M. C.

REGISTRATION NUMBER: 33,544

REFERENCE/DOCKET NUMBER: GI 5:

TELECOMMUNICATION INFORMATION:

TELEPHONE: (617) 876-1170

TELEPAX: (617) 876-5851

TELEPAX: (617) 876-5851

TELEPAX: (617) 876-5851
                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Vei
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/810,436
FILING DATE: 04-MAR-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
APPLICANT: Lavallie, Edward R.
TITLE OF INVENTION: THIOREDOXIN, THIOR
TITLE OF INVENTION: THIOREDOXIN-LIKE M
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TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 22
                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE,DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: UFILING DATE: 04-MAR-1CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/165,301
FILING DATE: 10-DEC-1993
ATTORNEY/AGENT INFORMATION:
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LENGTH: 109 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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STATE: Massachusetts
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6143524
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CHARACTERISTICS:
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04-MAR-1997
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PEPTIDE AND PROTEIN FUSIONS TO
THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
THIOREDOXIN-LIKE MOLECULES
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; LENGTH: 109 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-810-436-26
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片
                                                                                                          ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-810-436-28
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Best Local Similarity 100.
Matches 4; Conservative
                                                     Matches
                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                 TELEPHONE: (617) 876-11
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 29
CORRESSONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc.
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE:
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CITY: Cambridge
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33 CGPC 36
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                         2 CGPC 5
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                                                  4; Conservative
                                                                                                                                                 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Massachusetts
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LaVallie, Edward R.
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04-MAR-1997
V. ,,
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                                                                  100.0%; Score 31; DB 2;
100.0%; Pred. No. 3.5e+02;
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100.0%; Pred. No. 3.5e+02;
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                                                      Mismatches
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                                                                               Length 109
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                                                      Indels
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RESULT 55 US-09-485-885-17

Sequence 17, Application US/09485885 Patent No. 6342224

GENERAL INFORMATION:

Bruck, Claudine

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US-09-248-796A-19911
; Sequence 19911, Application US/09248796A
; Patent No. 6747137
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                                                                   RESULT 57
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SEQ ID NO 17
LENGTH: 109
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APPLICANT: CHEN, DAVID CI
APPLICANT: HU, NIEN-TAI
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Best Local Similarity
Matches 4; Conserv
GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                    OTHER INFORMATION: amino acid of thioredoxin -09-166-966E-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: CHEN, YUN-JU
APPLICANT: HSEU, TZONG-HSIUNG
TITLE OF INVENTION: TZONG-HSIUNG
FILE REFERENCE 32350-150960
CURRENT APPLICATION NUMBER: US/09/166,966E
CURRENT FILING DATE: 1998-10-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Lombardo-Bencheikh, Angela TITLE OF INVENTION: Vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: TW 86114750 PRIOR FILING DATE: 1997-10-08
                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 11
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                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 10
TYPE: PRT
                                                                                                                                                                                                                                                                                                                       PEATURE:
NAME/KEY: thioredoxin
                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                        LOCATION: 1..109
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Delisse, Anne-Marie Eva Fernande
Gerard, Catherine Marie Ghislaine
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                                                                                                                                                                                           100.0%; Score 31; DB 2; 100.0%; Pred. No. 3.5e+02; tive 0; Mismatches 0;
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100.0%; Pred. No. 3.5e+02;
cive 0; Mismatches 0;
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RESULT 58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCT-US94-14179-22
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Matches
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SEQ ID NO 19911
LENGTH: 109
TYPE: PRT
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 Matches
                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
                                                                                                                                                                                TELEFAX: (617) 876-58
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14179
                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                 REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy
                                                                                      MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: McCoy, John APPLICANT: DiBlasio-Sm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
y Match 100.0%; Score 31; DB 4; Length 109; Local Similarity 100.0%; Pred. No. 3.5e+02; hes 4; Conservative 0; Mismatches 0; Indels
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                                                                                                           TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                              NAME: Meinert, M.
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nilarity 100.0%;
Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genetics Institute,
                                                                                                                                                                            7) 876-5851
7 NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kathleen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PEPTIDE AND PROTEIN FUSIONS TO THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Edward R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>.</u>.
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Pred. No. 3.5e+02;
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RESULT 60
PCT-US94-14179-28
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; Sequence 26, Application PC/TUS9414179
; GENERAL INFORMATION:
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MOLECULE TYPE: protein
PCT-US94-14179-26
                                                                                                 Sequence 28, Application PC/TUS9414179

GENERAL INFORMATION:
APPLICANT: McCoy, John
APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 4; Conservative
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ver.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14179
FILING DATE:
CTROST-----
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APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy
                                                         NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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CORRESPONDENCE ADDRESS:
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CITY: Cambridge
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: GI 5188D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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                                       ADDRESSEE:
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Cambridge
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87 CambridgePark Drive
                 E: Genetics Institute, Inc
87 CambridgePark Drive
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(617) 876-5851
70 NO: 26:
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NUMBER OF SEQ ID NOS: 19335
SOPTMARE: Patent.pm
SEQ ID NO 4028
LENGTH: 110
TYPE: PAT
ORGANIAM: Homo sapiens
FEATURE:
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                                                                                                                         ; NAME/KEY: SIGNAL ; LOCATION: -102..-1 US-09-621-976-4028
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Patent No. 6639063
GENERAL INFORMATION:
                                                            Matches
                                                                      Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (617) 876-11:
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: ESTS and FILE REFERENCE: GENSET.054PR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Meinert, M. C.
REGIZENTON NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 0214v
COMPUTER READABLE FORM:
TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION: NAME: Meinert, M. C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: PCT/US94/14179
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74 CGPC
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                               2 CGPC 5
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 77
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100.0%; Pred. No. 3.5e+02;
ative 0; Mismatches 0;
                                                                           100.0%;
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                                                                              Score 31; Pred. No.
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                                                              Mismatches
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                                                                              3.5e+02;
                                                                                             DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 109
                                                                                          Length 110;
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                                                              Indels
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                                                              Gaps
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RESULT 62 US-09-264-419C-6 ; Sequence 6, Application US/09264419C ; Patent No. 6174682

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GENERAL INFORMATION:

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; OTHER INFORMATION: Description of Artificial Sequence: consensus ; OTHER INFORMATION: sequence US-09-264-419C-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 5674
SEQ ID NO 3859
LENGTH: 111
                                                                                                                                                                                            Sequence 6739, Application US/09107532A
Patent No. 6583275
GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conser
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Best Local Similarity
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SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Patent No. 6380370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/134,001C CURRENT FILING DATE: 1998-08-13 PRIOR APPLICATION NUMBER: US 60/064,964 PRIOR FILING DATE: 1997-11-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: MNI-076
CURRENT APPLICATION NUMBER: US/09/264,419C
CURRENT FILING DATE: 1999-03-08
NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 1997-08-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: LYNIN DOUGETTE-Stamm et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS FILE REFERENCE: GTC-007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Khodadous TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH: 111
                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                         APPLICANT: Lynn A Doucette-Stamm
TITLE OF INVENTION: NUCLEIC ACID
ENTEROCOCCUS
                                                                                                                                                                                                                                                                                                                            36 CGPC 39
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STATE: Massachusetts
COUNTRY: USA
ZIP: 02354
                                                    CITY: Waltham
                                                                     STREET:
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/ENTION: No. 6174682el Thioredoxin Family Active Site Molecules and
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                                                                 100 Beaver Street
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Pred. No. 3.5e+02;
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                                                                                                               and David Bush
AND AMINO ACID SEQUENCES RELATING TO
FARCIUM FOR DIAGNOSTICS AND THERAPEUTICS
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Sequence 3195, Application US/10104047

Patent No. 6943241

GENERAL INFORMATION:
APPLICANT: HELLX RESEARCH INSTITUTE
TITLE OF INVENTION: No. 6943241e1 full length cDN
FILE REFERENCE: H1-A0105

CURRENT APPLICATION NUMBER: US/10/104,047

CURRENT FILING DATE: 2002-03-25

PRIOR APPLICATION NUMBER:
PRIOR FILING DATE:
NUMBER: PRIOR FILING DATE:
NUMBER: PRIOR FILING DATE:
NUMBER OF SEQ ID NOS: 4096
SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 3199

LENGTH: 113
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US-10-104-047-3199
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                                                                                                                                                          ; ORGANISM: Homo sapiens US-10-104-047-3199
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                                                                            Matches
                                                                                               Query Match
Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                  TYPE: PRT
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APPLICATION NUMBER: US/09/107,532A

FILING DATE: 30-Jun-1998

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/085,598

FILING DATE: 14 May 1998

APPLICATION NUMBER: 60/051571

FILING DATE: July 2, 1997

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
COMPUTER: PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature LOCATION: (B) LOCATION 1...1 SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: protein HYPOTHETICAL: YES ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
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37 CGPC 40
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                                       2 CGPC 5
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SOFTWARE: ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Ariniello, Pamela Deneke REGISTRATION NUMBER: 40,489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Enterococcus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
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                                                                            100.0%; Score 31; DB 2;
100.0%; Pred. No. 3.6e+02;
tive 0; Mismatches 0;
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Pred. No. 3.5e+02;
); Mismatches 0;
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RESULT 66
US-08-918-288-71
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US-09-282-357-71
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No. 6238890
                                                                                                                                                                         Patent No. 6242580
GENERAL INFORMATION:
                                                                                                                                                                                             Sequence 71, Application US/09282357 Patent No. 6242580
                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local
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APPLICANT: BOIME, Irving
APPLICANT: MOYLE, William R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 114 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 08/853,524
APPLICATION NUMBER: 08/853,524
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: 08/199,382
FILING DATE: 18-FEB-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Murashige, Kate H
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29,
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-887-1500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: DOS
SOFTWARE: PASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/918,288
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatik
OPERATING SYSTEM: DOS
                                              NUMBER OF SEQUENCES: 83
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
                                                                                                  APPLICANT: BOIME, ITVING
APPLICANT: MOYLE, WILLIAM R.
TITLE OF INVENTION: SINGLE-CHAIN FORMS OF THE
TITLE OF INVENTION: GLYCOPROTEIN HORMONE QUARTET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
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CITY: Washington
STREET: Zuvu
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TOPOLOGY: lin
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CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                       90 CGPC 93
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                             2000 Pennsylvania Avenue, NW, suite 5500
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100.0%; P
tive 0;
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Pred. No. 3.6e+02;
); Mismatches 0;
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; ORGANISM: Arabidopsis thaliana US-09-897-425-41
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                                                                                                                                      PRIOR APPLICATION NUMBER: 09/210,843
PRIOR FILLING DATE: 1998-12-18
PRIOR FILLING DATE: 1998-12-18
PRIOR PELICATION NUMBER: 08/46,021
PRIOR FILLING DATE: 1997-04-25
PRIOR APPLICATION NUMBER: 08/366,783
PRIOR FILLING DATE: 1994-12-30
PRIOR FILLING DATE: 1994-12-30
PRIOR FILLING DATE: 1993-11-16
PRIOR APPLICATION NUMBER: 07/659,835
PRIOR APPLICATION NUMBER: 07/659,835
PRIOR FILLING DATE: 1991-02-22
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Best Local Similarity
Matches 4; Conserv
                                                                                SOFTWARE: Pa
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Patent No. 6750046
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: MOLONEY, MAURICE M.
APPLICANT: DALMIA, BIPIN K.
TITLE OF INVENTION: PREPARATION OF THIOREDOXIN AND THIOREDOXIN REDUCTASE
TITLE OF INVENTION: PROTEINS ON OIL BODIES
FILE REFERENCE: 034547/0106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 2001-07-03
                                                                                                                         NUMBER OF SEQ ID NOS: 55
                                     TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Murashige Kate H
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-887-1500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 536
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 08/9
FILLING DATE: 25 AUG-1997
APPLICATION NUMBER: 08/8
FILLING DATE: 08/8
                                                               LENGIH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 114 amino acid
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SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 09-MAY-1997
APPLICATION NUMBER: 08/199,382
FILING DATE: 18-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 202-887-0763
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/09/282,357
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100.0%; Pred. No. 3.6e+02;
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NAME/KEY: misc feature;
LOCATION: (B) LOCATION 1...114
SEQUENCE DESCRIPTION: SEQ ID NO: 2930:
US-09-107-433-2930
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RESULT 70
US-09-786-715-11
; Sequence 11, Application US/09786715
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Best Local Similarity luv.
Thes 4; Conservative
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Patent No. 6800744
Patent No. 6800744
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 2930:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/085131
PILING DATE: May 12, 1998
APPLICATION NUMBER: 60/05153
PILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-011
TELECHONE: (781)893-5007
TELECHONE: (781)893-8077
TELECAX: (781)893-8277
                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: protein
HYPOTHETICAL: YES
ORIGINAL SOURCE:
ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/107,433 FILING DATE: 30-Jun-1998 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                  FEATURE:
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MEDIUM TYPE: CD/ROM ISO9660
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ADDRESSEE: GENOME
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                                                                                            38 CGPC 41
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                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 114 amino acids TYPE: amino acid
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OPERATING SYSTEM: <Unknown>
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Pred. No. 3.6e+02;
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; Patent No. 6897356;
GENERAL INFORMATION:
APPLICANT: Allen, Steve
APPLICANT: Lu, Albert
APPLICANT: Thorpe, Cathy
TITLE OF INVENTION: Thioredoxin H Homologs
FILE REFERENCE: BB-1246
CURRENT APPLICATION NUMBER: US/09/786,715;
CURRENT APPLICATION NUMBER: 60/09/786,715
PRIOR APPLICATION NUMBER: 60/099,501
PRIOR FILING DATE: September 8, 1998
NUMBER OF SEQ ID NOS: 13
                                                                                                                                                ; SOFTWARE: Microsoft Office 97
; SEQ ID NO 11
; LENGTH: 114
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
US-09-786-715-11
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                                                                                            Query Match 100.0%;
Best Local Similarity 100.0%;
                                                                         Matches
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                                     2 CGPC 5
                                                                         4; Conservative
CGPC 43
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Search completed: February 23, 2006, 00:43:45 Job time: 44.5 secs

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Maximum DB
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/cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
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US-10-20-118A-3
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US-10-963-903-98
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US-10-293-157-20
US-10-424-599-255327
US-10-104-440-11
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72 31 100.0 345 4 US-10-255-567A-619 Sequence 73 31 100.0 345 4 US-10-467-252-8 Sequence 74 31 100.0 345 4 US-10-600-816-4 Sequence 75 31 100.0 345 5 US-10-500-428-2 Sequence	58 31 100.0 342 4 US-10-244-830-332 Sequence 69 31 100.0 342 4 US-10-427-717-332 Sequence 70 31 100.0 342 4 US-10-648-780-332 Sequence 71 31 100.0 344 4 US-10-757-701-38850 Sequence	66 31 100.0 342 4 US-10-195-835-332 Sequence 67 31 100.0 342 4 US-10-286-333-332 Sequence 69 31 100.0 342 4 US-10-286-333-332 Sequence	64 31 100.0 342 4 US-10-125-635A-332 Sequence 65 31 100.0 342 4 US-10-002-603-332 Sequence	62 31 100.0 342 3 US-09-938-864-332 Sequence 63 31 100.0 342 3 US-09-785-019-332 Sequence	61 31 100.0 341 5 US-10-948-518-158 Sequence	59	58 31 100.0 334 6 US-11-097-143-2163 Sequence	56 31 100.0 332 5 US-10-631-467-1495 Sequence	55 31 100.0 332 4 US-10-839-882-39 Sequence	53 31 100.0 331 5 US-10-450-763-36089 Sequence	52 31 100.0 331 4 US-10-017-161-618 Sequence	50 31 100.0 330 5 US-10-7/4-3584-2528 Sequence	49 31 100.0 330 5 US-10-774-355A-2525 Seguence	48 31 100.0 330 5 US-10-774-355A-2471 Sequence	47 31 100.0 330 5 US-10-774-355A-1805 Sequence	45 31 100.0 330 4 US-10-437-963-154386 Sequence	44 31 100.0 329 5 US-10-732-923-9551 Sequence	42 31 100.0 327 4 US-10-032-585-7827 Segmence	31 100.0 326 4 US-10-264-049-3112 Sequence	39	38 31 100.0 325 5 US-10-357-819-28 Sequence	36	35 31 100.0 325 4 US-10-315-664-74 Sequence	33 31 100.0 325 3 US-09-978-360A-406 Sequence	31 100.0 325 3 US-10-902-31-1/ sequence	30 31 100.0 323 4 US-10-343-650A-2 Sequence	28 31 100.0 323 4 US-10-024-212-106 Sequence	27 31 100.0 323 4 US-10-141-531-28 Sequence	26 31 100.0 322 4 US-10-292-798-542 Sequence	24 31 100.0 321 4 US-10-282-122A-75634 Segmence	23 31 100.0 321 4 US-10-041-615-71 Sequence	21 31 100.0 321 4 US-10-024-399-30 sequence	20 31 100.0 321 4 US-10-024-399-28 Sequence	18	17 31 100.0 321 4 US-10-023-601-48 Sequence	16 31 100.0 321 4 US-U9-771-730-43 sequence	14 31 100.0 320 5 US-10-774-355A-2225 Sequence	13 31 100.0 318 5 US-10-732-923-9556 Sequence	12 31 100.0 318 4 US-10-437-963-132260 Sequence	10 31 100.0 317 4 US-10-276-774-2644 Sequence	09 31 100.0 314 4 US-10-479-284-16 Sequence	07 : 31 100.0 309 4 US-10-369-493-19683 Sequence	06 31 100.0 306 4 US-10-267-430-13 Sequence	05 31 100.0 304 5 US-10-902-531-16 Sequence
72 31 100.0 345 4 US-10-252-567A-619 Sequence 8, 73 31 100.0 345 4 US-10-467-252-8 Sequence 8, 74 31 100.0 345 4 US-10-600-816-4 Sequence 4, 75 31 100.0 345 5 US-10-500-428-2 Sequence 2,	58 31 100.0 342 4 US-10-244-830-332 69 31 100.0 342 4 US-10-427-717-332 70 31 100.0 342 4 US-10-648-780-332 71 31 100.0 344 4 US-10-767-701-39850	66 31 100.0 342 4 US-10-195-835-332 Sequence 33: 67 31 100.0 342 4 US-10-286-333-332 Sequence 33: 67 31 100.0 342 4 US-10-195-835-332 S	64 31 100.0 342 4 US-10-125-635A-332 Sequence 33:	62 31 100.0 342 3 US-09-938-864-332 Sequence 33: 63 31 100.0 342 3 US-09-785-019-332 Sequence 33:	61 31 100.0 341 5 US-10-948-518-158 Sequence 15	59 31 100.0 341 4 US-10-106-698-4514 Sequence 45:	58 31 100.0 334 6 US-11-097-143-2163 Sequence 21	56 31 100.0 332 5 US-10-631-467-1495 Sequence	55 31 100.0 332 4 US-10-839-882-39 Sequence	53 31 100.0 331 5 US-10-450-763-36089 Sequence	52 31 100.0 331 4 US-10-017-161-618 Sequence	50 31 100.0 330 5 US-10-7/4-3584-2528 Sequence	49 31 100.0 330 5 US-10-774-355A-2525 Seguence	48 31 100.0 330 5 US-10-774-355A-2471 Sequence	47 31 100.0 330 5 US-10-774-355A-1805 Sequence	45 31 100.0 330 4 US-10-437-963-154386 Sequence	44 31 100.0 329 5 US-10-732-923-9551 Sequence	42 31 100.0 327 4 US-10-032-585-7827 Segmence	31 100.0 326 4 US-10-264-049-3112	39	38 31 100.0 325 5 US-10-357-819-28 Sequence 28	36	35 31 100.0 325 4 US-10-315-664-74 Sequence 74	33 31 100.0 325 3 US-09-978-360A-406 Sequence 40	31 100.0 325 3 US-09-800-095A-96 Sequence 1/	30 31 100.0 323 4 US-10-343-650A-2 Sequence 2,	28 31 100.0 323 4 US-10-024-212-106 Sequence 20	27 31 100.0 323 4 US-10-141-531-28 Sequence 28	26 31 100.0 322 4 US-10-292-798-542 Sequence 54	24 31 100.0 321 4 US-10-282-122A-75634 Seguence 756 25 31 100.0 321 5 US-10-774-355A-1807 Seguence 180	23 31 100.0 321 4 US-10-041-615-71 Sequence 71,	21	20 31 100.0 321 4 US-10-024-399-28 Sequence 28,	18	17 31 100.0 321 4 US-10-023-601-48 Sequence 48,	15	14 31 100.0 320 5 US-10-774-355A-2225 Sequence 222	13 31 100.0 318 5 US-10-732-923-9556 Sequence 955	12 31 100.0 318 4 US-U9-771-730-20 Sequence 20,	10 31 100.0 317 4 US-10-276-774-2644 Sequence 264	09 31 100.0 314 4 US-10-479-284-16 Sequence 16,	07 31 100.0 309 4 US-10-369-493-19683 Sequence 196	06 31 100.0 306 4 US-10-267-430-13 Sequence 13,	05

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APPLICANT: McBride, Jere W.
APPLICANT: Walker, David H.
TITLE OF INVENTION: Ehrlichia Disulfide Bond Formation
TITLE OF INVENTION: Proteins and Uses Thereof
FILE REFERENCE: D6428
CURRENT APPLICATION NUMBER: US/10/286,516A
CURRENT FILING DATE: 2002-11-01
PRIOR APPLICATION NUMBER: US 60/335,611
PRIOR APPLICATION NUMBER: US 60/335,611
PRIOR PILING DATE: 2001-11-01
NUMBER OF SEQ ID NOS: 20
SEQ ID NO 11
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US-10-286-516A-11
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            Sequence 1, Application US/10234319A
Publication No. US20030109457A1
GENERAL INFORMATION:
APPLICANT: Atlas, Daphne
TITLE OF INVENTION: MULTI-COMPONENT ANTIOXIDANT COMPOUNDS, PHARMACBUTICAL COMPOSITION
TITLE OF INVENTION: CONTAINING SAME AND THEIR USE FOR REDUCING OR PREVENTING OXIDATI
FILE REFERENCE: 24890
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Best Local
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                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Escherichia coli
PEATURE:
OTHER INFORMATION: Escherichia coli conserved cysteine
OTHER INFORMATION: motif in active site of thioredoxin
OTHER INFORMATION: protein
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US-09-804-291-441
US-09-804-291-441
US-10-343-650A-448
US-10-316-441
US-10-819-312-58
US-10-369-493-21120
US-10-369-493-21120
US-10-10-294-025-1011
US-10-369-493-215-1011
US-10-369-493-215-1011
US-10-369-493-215-1011
US-10-369-493-215-1011
US-10-369-493-215-1011
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US-10-369-493-215-101
US-10-369-493-215-101
US-10-435-596-52
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US-10-425-115-326082

US-10-928-992-162

US-09-878-178-2235

US-09-878-178-2235

US-10-046-935-2235

US-10-066-543-1422
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US-10-902-531-18
                                                                                                                                                                                                                                              Score 31; DB 4; Length 4; Pred. No. 1.7e+06; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 181709,
Sequence 441, Appl
Sequence 441, Appl
Sequence 441, Appl
Sequence 441, Appl
Sequence 297, Appl
Sequence 21120, A
Sequence 21120, A
Sequence 1011, Ap
Sequence 1011, Ap
Sequence 1011, Ap
Sequence 1011, Ap
Sequence 41715, A
Sequence 41785, A
Sequence 41585, A
Sequence 41585, A
Sequence 52, Appl
Sequence 162, Appl
Sequence 2235, Ap
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SOFTWARE: Patentin version 3.1
SEQ ID NO 1
LENGTH: 4
TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
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                                                                                                                                                                    RESULT 4
US-10-660-118A-1
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US-10-369-094-1
         GENERAL INFORMATION:
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum FILE REFERENCE: 2879-98
FILE REFERENCE: 2879-98
CURRENT APPLICATION NUMBER: US/10/660,118A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/499,960
PRIOR APPLICATION NUMBER: 60/499,960
PRIOR FILING DATE: 2002-09-10
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
                                                                                                                                   Sequence 1, Application US/10660118A
Publication No. US20040131606A1
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Matches 4; Conserv
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APPLICANT: Marks, Paul A.
APPLICANT: Rifkind, Richard A.
APPLICANT: Butler, Lisa M.
TITLE OF INVENTION: Method of Treating TRX Mediated
FILE REFERENCE: 3254.1005.001
                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 4
TYPE: PRT
ORGANISM: homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/369,094
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: U.S. 60/357,383
PRIOR FILING DATE: 2002-02-15
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PRIOR APPLICATION NUMBER: 60/462,082
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LOCATION: (1)...(1)
OTHER INFORMATION: ACETYLATION
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100.0%; Pred. No.
tive 0; Mismatch
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/628,391
CURRENT FILING DATE: 2003-07-29
PRIOR APPLICATION NUMBER: PCT/IE02/00011
PRIOR FILING DATE: 2002-01-30
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.2
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/10628391 Publication No. US20040146526A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                   Sequence 149, Application US/10776933
Publication No. US20040241717A1
                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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Best Local
                                                             APPLICANT: HAMSEN, BO
APPLICANT: THRUE, CHARLOTTE ALBAEK
APPLICANT: WESTERGAARD, MAJKEN
APPLICANT: WESTERGAARD, MAJKEN
APPLICANT: WESTERGARD, MARGIT
TITLE OF INVENTION: OLICOMERIC COMPOUNDS FOR THE MODULATION OF THIOREDOXIN
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: 58614 (71432)
CURRENT APPLICATION NUMBER: US/10/776,933
CURRENT FILING DATE: 2004-02-10
CURRENT FILING DATE: 2004-02-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: KELLEHOR, Dermot
APPLICANT: ABDEL-LATIF, MOHAMED M. TITLE OF INVENTION: INHIBITION OF NF-KappaB ACTIVATION
FILE REFERENCE: P69048USO
                                 PRIOR APPLICATION NUMBER: 60/446,374
PRIOR FILING DATE: 2003-02-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: WINDLE, Henry J. APPLICANT: O'TOOLE, Dermot
                   NUMBER OF SEQ ID NOS: 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
OTHER INFORMATION: the redox active peptide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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PatentIn Ver. 3.2
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Pred. No. 1
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APPLICANT: Kwon, Mijung
TITLE OF INVENTION: Anti-angiogenesis methods, con
FILE REFERENCE: ME02-001
CURRENT APPLICATION NUMBER: US/10/304,287
CURRENT FILING DATE: 2002-11-26
PRIOR APPLICATION NUMBER: US 60/333,866
PRIOR APPLICATION NUMBER: US 60/333,866
PRIOR FILING DATE: 2001-11-28
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Microsoft Word
SEQ ID NO 4
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 233, A Publication No.
APPLICANT: Luginbuhl, Peter
APPLICANT: Muchhal, Umesh
TITLE OF INVENTION: Nucleic Acids and Proteins with Thioredoxin Reductase Activity
FILE REFERENCE: A-71457-3
CURRENT FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: US 60/370,609
PRIOR FILING DATE: 2002-04-05
PRIOR APPLICATION NUMBER: US 60/376,682
PRIOR APPLICATION NUMBER: US 60/376,682
PRIOR PILING DATE: 2002-04-29
PRIOR PILING DATE: 2002-05-06
PRIOR APPLICATION NUMBER: US 60/289,029
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 239
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APPLICANT: Dalmia, Bipin
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                                                                                                                                                                                                                                                                                                     Heifetz, Peter
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Pred. No. 1.7e+06;
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Publication No. US20040019104A1
GENERAL INFORMATION:
APPLICANT: Ebright, Richard H.
APPLICANT: Ebright, Yon W.
TITLE OF INVENTION: Reagents and Procedures
FILE REFERENCE: 744-34
           Sequence 2, Application US/10660118A

Publication No. US20040131606A1

GENERAL INFORMATION:

APPLICANT: White, Carl W.

TITLE OF INVENTION: Product and Process for Liquefaction of Mucus of Title Reference: 2879-98

CURRENT APPLICATION NUMBER: US/10/660,118A

CURRENT APPLICATION NUMBER: 00/409,960

PRIOR APPLICATION NUMBER: 60/409,960

PRIOR PILING DATE: 2002-09-10

PRIOR FILING DATE: 2002-09-10
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Matches 4; Conserv
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Best Local Similarity
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CURRENT FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: US 60/388,699
PRIOR FILING DATE: 2002-06-14
NUMBER OF SEQ ID NOS: 11
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EQ ID NO 233
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NAME/KEY: MISC FEATURE
1...(1)...(1)
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PRIOR APPLICATION NUMBER: 60/462,082
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ORGANISM: Artificial Sequence
FEATURE:
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OTHER INFORMATION: "Xaa" at position
OTHER INFORMATION: hydrophobic amino
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Pred. No.
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US-10-735-577-4
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US-10-660-118A-3
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                                                                   GENERAL INFORMATION:
APPLICANT: Waisman, David M.
TITLE OF INVENTION: Compositions and Methods
FILE REFERENCE: ME03-009
CURRENT FILING DATE: 2003-12-12
PRIOR APPLICATION NUMBER: US/10/735,577
PRIOR FILING DATE: 2002-12-13
PRIOR FILING DATE: 2002-12-13
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NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.1
SEQ ID NO 2
LENGTH: 6
TYPE: PRT
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Publication No. US20040142897A1
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 31; DB 4; Best Local Similarity 100.0%; Pred. No. 1.7e+06; Matches 4; Conservative 0; Mismatches 0;
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SOFTWARE: Microsoft Word
SEQ ID NO 4
LENGTH: 6
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CURRENT APPLICATION NUMBER: US/10/660,118A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR PPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR PILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin version 3.1
                                                   PRIOR FILING DATE: 2002-12-13
NUMBER OF SEQ ID NOS: 160
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NAME/KEY: misc feature
LOCATION: (1)...(6)
OTHER INFORMATION: Xaa = any amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 6
TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
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Pred. No.
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; OTHER INFORMATION: C
US-10-954-951-11
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US-10-461-224-10
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US-10-735-577-4
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PRIOR FILING DATE: 2002-06-14
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin version 3.2
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
 Query Match
Best Local Similarity
Matches 4; Conserv
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Matches 4; Conserv
                                                                                                                                                            SOFTWARE: PatentIn version 3.3 SEQ ID NO 11
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APPLICANT: Ebright, Richard H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10, Application US/10461224 Publication No. US20040019104A1
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TITLE OF INVENTION: Reagents and Procedures for High Specificity Labeling FILE REFERENCE: 1744-34

CURRENT APPLICATION NUMBER: US/10/461,224

CURRENT FILING DATE: 2003-06-13
                                                                                                                                                                                                                                                                       APPLICANT: Kudlicki, Wieslaw Antoni
APPLICANT: Keppetipola, Shiranthi
TITLE OF INVENTION: Compositions and Methods For Synthesizing,
TITLE OF INVENTION: Detecting Biomolecules
FILE REFERENCE: 0942.6650001
                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/508,142
PRIOR FILING DATE: 2003-10-01
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/954,951
CURRENT FILING DATE: 2004-10-01
                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Hanson, George
                                                                                                                                 TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
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                                                                                                             ORGANISM: Artificial
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Pred. No. 1.7e+06;
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 DB 5; I
1.7e+06;
les 0;
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RESULT 17
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                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Complementary peptide ligands from the human genome FILE REFERENCE: Human patent CURRENT APPLICATION NUMBER: US/09/572,404B CURRENT FILING DATE: 2000-05-17 NUMBER OF SEQ ID NOS: 4203 SOFTWARE: ProtPatent version 1.0 SEQ ID NO 970 LENGTH: 10 TYPE: PRT
                 Sequence 2, Application US/10615343
Publication No. US20050009110A1
GENERAL INFORMATION:
APPLICANT: CHANG, XIAO-JIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 970, Applic Publication No. US20 GENERAL INFORMATION:
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APPLICANT: Proteom Ltd
APPLICANT: Proteom Ltd
APPLICANT: Proteom Ltd
TITLE OF INVENTION: Complementary peptide ligands from the human genome
FILE REFERENCE: Human patent
CURRENT APPLICATION NUMBER: US/09/572,404B
CURRENT FILING DATE: 2000-05-17
NUMBER OF SEQ ID NOS: 4203
                                                                                                                                                                                                                                    Matches
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Best Local Similarity
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TITLE OF INVENTION: METHODS OF PRODUCING ANTIBODIES FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Proteom Ltd
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TYPE: PRT
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nilarity 100.0%;
Conservative 0
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Pred. No. 2.1e+02;
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Pred. No. 2.1e+02
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-615-343-2
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                                                                                                                                             US-10-044-844-217
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CURRENT APPLICATION NUMBER: US/10/615,343
CURRENT FILING DATE: 2003-07-08
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
                                                                             Query Match
Best Local Similarity
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Best Local (
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                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 25231-20061.03
TELECOMPUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
TELEPHONE: (650) 494-0792
TELEPAX: (650) 494-0792
INFORMATION FOR SEQ ID NO: 217:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Y Match 100.0%;
Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/044,844
FILING DATE: 10-Jan-2002
CLASSIFICATION: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: CATHERINE M. POLIZZI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 217:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER LLP
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: APL IMMUNOREACTIVE PEPTIDES, CONJUGATES THEREOF AND METHODS OF TREATMENT FOR APL ANTIBODY-MEDIATED PATHOLOGIES
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Marquis, David M.
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 CGPC
                                   CGPC 5
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TYPE: amino acid
STRANDEDNESS: single
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                                                                       Conservative
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                                                                                    100.0%;
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                                                                     Score 31; DB 4; : Pred. No. 2.9e+02; ); Mismatches 0;
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Pred. No. 2.6e+02;
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Sequence 10, Application US/09963339

Publication No. US20030049700A1

GENERAL INFORMATION:
APPLICANT: Bandaru, Rajasekhar
TITLE OF INVENTION: 22108 AND 47916, NOVEL HUMAN THIOREDOXIN
TITLE OF INVENTION: FAMILY MEMBERS AND USES THEREOF
FILE REFERENCE: 10448-090001
CURRENT APPLICATION UMMBER: US/09/963,339
CURRENT FILING DATE: 2001-09-25
PRIOR APPLICATION NUMBER: 60/235,049
PRIOR FILING DATE: 2000-09-25
INUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 4.0
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GENERAL INFORMATION:
APPLICANT: Victo
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 217:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/846,079
FILING DATE: 13-May-2004
CLASSIFICATION: CUNknown>
ATTORNEY/ACENT INFORMATION:
NAME: CATHERINE M. POLIZZI
REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 25231-20061.03
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 217:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 226 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yu, Lin
TITLE OF INVENTION: APL IMMUNOREACTIVE PEPTIDES, CONJUGATES THEREOF AND
METHODS OF TREATMENT FOR APL ANTIBODY-MEDIATED PATHOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 CGPC 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Conservative
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TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: MORRISON & FOERSTER LLP
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o. US20050208480A1
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Marquis, David M.
Jones, David S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 31; DB 5;
100.0%; Pred. No. 2.9e+02;
tive 0; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/10/050,882
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: 09/661,453
PRIOR FILING DATE: 2000-09-13
PRIOR APPLICATION NUMBER: PCT/US00/06783
PRIOR APPLICATION NUMBER: PCT/US00/06783
PRIOR FILING DATE: 2000-03-16
PRIOR APPLICATION NUMBER: 60/125,055
PRIOR FILING DATE: 1999-03-18
NUMBER OF SEQ ID NOS: 156
SOPTWARE: Patentin Ver. 2.0
SEQ ID NO 98
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LENGTH: 25
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-050-882-98
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; ORGANISM: Homo sapiens
US-10-145-586-59
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SEQ ID NO 59
LENGTH: 19
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Best Local Similarity
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Best Local Similarity
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CURRENT FILING DATE: 2002-05-14
Prior Application removed - See File Wrap;
NUMBER OF SEQ ID NOS: 95
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APPLICANT: Kapeller-Libermann, Rosana
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS,
TITLE OF INVENTION: HUMAN THIOREDOXIN FAMILY MEMBERS, HUMAN LEUCINE-RICH
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY MEMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Alexandra Glucksmann, Maria
APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: M. Galvin, Katherine
APPLICANT: Weich, Nadine
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TYPE: PRT
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100.0%; Pred. No. 3.3e+02;
vative 0; Mismatches 0;
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PRIOR PELICATION NUMBER: US 10/050,882
PRIOR PELICATION NUMBER: US 09/661,453
PRIOR PELICATION NUMBER: US 09/661,453
PRIOR PELICATION NUMBER: PCT/US00/06783
PRIOR PELICATION NUMBER: PCT/US00/06783
PRIOR PELICATION NUMBER: US 60/125,055
PRIOR PELICATION
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US-10-179-382-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 31, Application US/10179382
Publication No. US20030166235A1
GENERAL INFORMATION:
APPLICANT: THIREOS, GEORGE
APPLICANT: KAFETZOPOULOS, DIMITRIS
TITLE OF INVENTION: DNA ENCODING AN ARTHROPOD CHITIN SYNTHASE
FILE REFERENCE: DNA ENCODING AN ARTHROPOD CHITIN SYNTH
CURRENT FILING DATE: 2002-06-26
PRIOR APPLICATION NUMBER: US/10/179,382
CURRENT FILING DATE: 2002-06-26
PRIOR APPLICATION NUMBER: US/09/230,041
PRIOR FILING DATE: 1999-01-19
                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 31
LENGTH: 28
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                                                                                                                                       Query Match
Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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21 CGPC 24
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                                                                                                                                       100.0%; Score 31; DB 4;
100.0%; Pred. No. 4.2e+02;
tive 0; Mismatches 0;
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100.0%; Pred. No. 3.9e+02;
tive 0; Mismatches 0;
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100.0%; Pred. No. 3.9e+02;
tive 0; Mismatches 0;
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FOR

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APPLICANT: Rupp, Keith;
APPLICANT: Bupp, Keith;
APPLICANT: University of Medicine and Dentistry of New Jersey
ITITLE OF INVENTION: Targeting Viral Vectors to Specific Cells
FILE REFERENCE: 601-1-095
CURRENT APPLICATION NUMBER: US/09/881,572A
CURRENT FILING DATE: 2002-03-05
PRIOR APPLICATION NUMBER: 60/212,239
PRIOR APPLICATION NUMBER: 60/212,239
PRIOR FILING DATE: 2000-06-16
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn version 3.0
SEQ ID NO 5
LENGTH: 30
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; ORGANISM: Amphotropic Murine Leukemia Virus
US-09-881-572A-25
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US-09-881-572A-5
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US-09-864-761-40069
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 25
Sequence 40069, Application US/09864761
Patent No. US20020048763A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
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Best Local Similarity 100.0%;
Matches 4; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 25, Application US/09881572A
Patent No. US20020164583A1
                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/881,572A CURRENT FILING DATE: 2002-03-05 PRIOR APPLICATION NUMBER: 60/212,239 PRIOR FILING DATE: 2000-06-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Bupp, Keith;
APPLICANT: University of Medicine and Dentistry of New Jersey
TITLE OF INVENTION: Targeting Viral Vectors to Specific Cells
FILE REFERENCE: 601-1-095
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                                                                                                                                                                                                                                                                                                                                                                                            murine leukemia virus
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Pred. No. 4.4e+02;
Mismatches 0;
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Pred. No.
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PRIOR FILING DATE: 2000-10-04
PRIOR PRICING NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-08-03
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SEQ ID NO 40069
                                                                                                                                          Query Match
Best Local
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NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence Listing Engine vers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00669
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-06-30 PRIOR APPLICATION NUMBER: US 09/774,203
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PRIOR APPLICATION NUMBER: PCT/US01/00663
                                                                                                                                                                                                                                                OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 5.4
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 4.6
OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 4.4
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 4.9
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 4.3
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 4.3
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 5
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 4
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 4
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 4
OTHER INFORMATION: SWISSPROT HIT: BF446535.1, EVALUE 6.00e-12
OTHER INFORMATION: SWISSPROT HIT: Q02597, EVALUE 1.70e+00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: MAP TO AC003681.1
OTHER INFORMATION: EXPRESSED IN LUNG
OTHER INFORMATION: EXPRESSED IN HELA
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TYPE: PRT
ORGANISM: Homo sapiens
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FILING DATE: 2001-01-30
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                                                                                                               4; Conservative
                                                                                                                                             Similarity
                                                         CGPC 5
   CGPC
11
                                                                                                            100.0%; So
100.0%; P
                                                                                                               Score 31; DB
Pred. No. 4.6
0; Mismatches
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RESULT 28 US-10-424-599-160512

Sequence 160512, Application US/10424599

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RESULT 30
US-09-801-260-5
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US-10-424-599-160512
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; ORGANISM: Androctonus mauretanicus
US-10-044-359-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: SCORPION TOXINS
FILE REFERENCE: BB1367 US M.
CURRENT APPLICATION NUMBER: US/10/044,359
CURRENT FILING DATE: 2002-01-11
PRIOR APPLICATION NUMBER: 09/599,416
PRIOR FILING DATE: 2000-06-22
PRIOR FILING DATE: 1999-06-22
PRIOR FILING DATE: 1999-06-22
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APPLICANT: Cao Yongwei
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 160512
Sequence 5, Application US/09801260
Patent No. US20020034801A1
GENERAL INFORMATION:
APPLICANT: Curtis, Rory A.J.
TITLE OF INVENTION: 22105, A NOVEL HUMAN THIOREDOXIN FAMILY
TITLE OF INVENTION: MEMBER AND USES THEREOF
FILE REFERENCE: 10448-022001
CURRENT APPLICATION NUMBER: US/09/801,260
CURRENT FILING DATE: 2001-03-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Herrman, Rafael
APPLICANT: Wong, James F.
APPLICANT: Lee, Jian-Ming
                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Microsoft Office 97
SEQ ID NO 26
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Glycine max
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/10044359
                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 31; DB 4; ilarity 100.0%; Pred. No. 4.9e+02; Conservative 0; Mismatches 0:
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US-10-145-586-44
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                                                                                                                                                                                                                                                US-10-450-763-51435
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                                                                                                                                     Sequence 51435, Application US/10450763
Publication No. US20050196754A1
GENERAL INFORMATION:
APPLICANT: Hyseq, Inc.
TITLE OF INCENTION: NOVEL NUCLEIC ACIDS AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 44, Application US/10145586 Publication No. US20030138890A1 GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-03-07
NUMBER OF SEQ ID NOS: 5
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 36
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 44
LENCTH: 36
TYPE: PRT
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        FILE REFERENCE: 790CIP3/US
CURRENT APPLICATION NUMBER: US/10/450,763
CURRENT FILING DATE: 2003-06-11
PRIOR APPLICATION NUMBER: PCT/US01/08631
PRIOR FILING DATE: 2001-03-30
PRIOR FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/540,217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/145,586
CURRENT FILING DATE: 2002-05-14
Prior Application removed - See File Wrapper or NUMBER OF SEQ ID NOS: 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Bandaru, Rajasekhar
APPLICANT: Kapeller-Libermann, Rosana
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS,
TITLE OF INVENTION: HUMAN THIOREDOXIN FAMILY MEMBERS, HUMAN LEUCINE-RICH
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY MEMBER
FILE REFERENCE: 10448-188001
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PRIOR FILING DATE: 2000-03-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: consensus sequence
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Similarity 100.0%; Pred. No. 5e-
4; Conservative 0; Mismatches
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Weich, Nadine
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US-09-864-761-35367; Sequence 35367, Application US/09864761; Patent No. US20020048763A1
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                                                                                                                               CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00669
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR APPLICATION NUMBER: PCT/US01/006670
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PRIOR PRIOR DATE: 2000-09-21
PRIOR PRIOR PRIOR PRIOR DATE: 2000-09-21
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PRIOR FILING DATE: 2000-08-23
NUMBER OF SEQ ID NOS: 60736
SOPTWARE: CUSTOM
SEQ ID NO 51435
LENGTH: 42
NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence
SEQ ID NO 35367
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
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APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                   PRIOR APPLICATION |
PRIOR FILING DATE:
                                                                                                                APPLICATION NUMBER: US 09/774,203
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l, David K.
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Pred. No.
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US-10-424-599-255327; Requence 255327, Application US/10424599; Publication No. US20040031072A1
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US-10-293-157-20
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Matches
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Publication No. US20030144200A1
GENERAL INFORMATION:
APPLICANT: BAIRD, ANDREW
APPLICANT: ANDREASON, GRAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 20
                                                                                                                                                 GENERAL INFORMATION:
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Best Local (
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CURRENT APPLICATION NUMBER: US/10/293,157
CURRENT FILING DATE: 2002-11-12
PRIOR APPLICATION NUMBER: US/09/244,583
PRIOR FILING DATE: 1999-02-04
PRIOR APPLICATION NUMBER: 60/073,979
PRIOR APPLICATION NUMBER: 60/073,979
               APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
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NUMBER OF SEQ ID NOS: 29
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TITLE OF INVENTION: VASCULAR ENDOTHELIAL CELL GROWTH FACTOR: VEGF
CURRENT APPLICATION NUMBER: US/10/424,599
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NN: EXPRESSED IN BRAIN, SIGNAL = 2.7

NN: EXPRESSED IN HOME WARROW, SIGNAL = 3.4

NN: EXPRESSED IN FETAL LIVER, SIGNAL = 3.4

NN: EXPRESSED IN ADULT LIVER, SIGNAL = 2.4

NN: EXPRESSED IN HOLLT LIVER, SIGNAL = 2.4

NN: EXPRESSED IN HELLA, SIGNAL = 2.2

NN: EXPRESSED IN HELLA, SIGNAL = 1.8

NN: EXPRESSED IN PLACENITA, SIGNAL = 3.2

NN: EXPRESSED IN HELART, SIGNAL = 3.2

NN: EXPRESSED IN HELART, SIGNAL = 3.0

NN: EXPRESSED IN HELART, SIGNAL = 3.00e-03

NN: EXPRESSED IN HELART, SIGNAL = 3.00e-03

NN: EXPRESSED IN HELART, SIGNAL = 3.00e-04
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Pred. No. 5.7
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NUMBER OF SEQ II SEQ ID NO 255327

ID NOS:

2003-04-28 285684

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LENGTH:

CURRENT FILING DATE:

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; OTHER INFORMATION: Clone ID: PAT_MRT3847_72582C.1.pep
US-10-424-599-255327
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                                                               CURRENT APPLICATION NUMBER: US/10/104,440
CURRENT FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: 09/580,803
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 60/069,155
PRIOR FILING DATE: 1997-12-09
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR FILING DATE: 1997-12-29
PRIOR APPLICATION NUMBER: 60/078,541
PRIOR APPLICATION NUMBER: 60/078,541
PRIOR APPLICATION NUMBER: 60/078,541
PRIOR FILING DATE: 1998-03-19
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Best Local Similarity 100.
""" hes 4; Conservative
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Best Local Similarity 100.
Warches 4; Conservative
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SEQ ID NO 11
LENGTH: 44
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                                                                                                                                                                                                                                                                                                                                                            Sequence 11, Application No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Kovesdi, Imre
APPLICANT: Kessler, Paul
TITLE OF INVENTION: VEGF FUSION PROTEINS
FILE REFERENCE: 205654
CURRENT APPLICATION NUMBER: US/09/832,355A
CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 126
                                 SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                 TITLE OF INVENTION: ANTAGONISTS OF NEUROPILIN RECEPTOR FUNCTION AND TITLE OF INVENTION: THEREOF FILE REFERENCE: 48802 C
                                                    NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                         APPLICANT: KLAGSBRUN, Michael
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TYPE: PRT
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ORGANISM: Glycine max
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MIAO, Hua Quan
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Pred. No. 5.7
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US-10-424-599-252212
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US-10-104-610-15
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CURRENT FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: 09/580,989
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 60/069,155
PRIOR FILING DATE: 1997-12-09
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR APPLICATION NUMBER: 60/069,687
PRIOR FILING DATE: 1997-12-12
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Best Local Similarity 100.
Best Local Similarity 200.
                                                                                                                                         NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 252212
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                                                                                                                                                                                                                                                                                 APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                             APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
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TITLE OF INVENTION: SOLUBLE INHIBITORS OF VASCULAR ENDOTHELIAL GROWTH
TITLE OF INVENTION: FACTOR
TITLE OF INVENTION: AND USE THEREOF
FILE REFERENCE: 48801 C
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TYPE: PRT
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                                                                                                     LENGTH: 45
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: human
                                      NAME/KEY: unsure
LOCATION: (1)..(45)
OTHER INFORMATION: unsure at
                                                                                ORGANISM: Glycine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 CGPC 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 31; DB 4; 100.0%; Pred. No. 5.8e+02;
  all Xaa locations
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 44;
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US-10-425-115-210616; Sequence 210616, Application US/10425115; Publication No. US20040214272A1
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                                                ; OTHER INFORMATION: Clone ID: MRT4577_56384C.1.pep US-10-425-115-321622
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Best Local Similarity 100.
"Arches 4; Conservative
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APPLICANT: Kovalic, David K.
APPLICANT: Chou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT EAPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 210616
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Best Local Similarity
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                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 321622
LENGTH: 47
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 321622, Application US/10425115 Publication No. US20040214272A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                 APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                               APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules :
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
                                                                             FEATURE: Unsure
LOCATION: (1)..(47)
OTHER INFORMATION: unsure at
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Clone ID: MRT4577_12367C.1.peg
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                                                                                                                                                                ORGANISM: Zea mays
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100.0%;
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                                                                                                   all Xaa locations
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Pred. No. 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 31; DB 4;
Pred. No. 6e+02;
 Score 31;
Pred. No.
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 DB 4;
6e+02;
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                                                                                                                                                                                                                                                                                                                    and Other Molecules Associated With
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                 Length 47;
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules an
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 334172
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US-10-425-115-334172
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FULRENT APPLICATION NUMBER: US/11/097,143

CURRENT APPLICATION NUMBER: 2005-04-04

PRIOR APPLICATION NUMBER: 60/157,832

PRIOR FILING DATE: 1999-10-05

PRIOR PPLICATION NUMBER: 60/160,191

PRIOR FILING DATE: 1999-10-19

PRIOR PRIOR DATE: 1999-10-19

PRIOR APPLICATION NUMBER: 60/161,932

PRIOR APPLICATION NUMBER: 60/164,769

PRIOR PILING DATE: 1999-10-28

PRIOR PILING DATE: 1999-10-28

PRIOR FILING DATE: 1999-10-28

PRIOR FILING DATE: 1999-10-28
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PRIOR FILING DATE: 1999-12-28
PRIOR APPLICATION NUMBER: 60/175,693
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/184,831
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 2000-03-23
PRIOR FILING DATE: 2000-03-23
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SEQ ID NO 37521
LENGTH: 47
TYPE: PRT
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Publication No. US20040214272A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conserv
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Publication No. US20050208558A1
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APPLICANT: et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: DESCRIPTION: DESCRIPTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 43008
LENGTH: 49
TYPE: PRT
ORGANISM: Zea mays
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Pred. No. 6e+02;
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FEATURE:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-801-260-4
                                            SEQ ID NO 43
LENGTH: 52
TYPE: PRT
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 52
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APPLICANT: Curtis, Rory A.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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Best Local Similarity
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/09801260 Patent No. US20020034801A1
                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bandaru, Rajasekhar
APPLICANT: Kapeller-Libermann, Rosana
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS
TITLE OF INVENTION: HUMAN THIOREDOXIN FAMILY MEMBERS, HUMAN LEGCIN
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY MEMBERS FAMILY FAMI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Alexandra Glucksmann, Maria APPLICANT: Silos-Santiago, Inmaculada
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                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/145,586
CURRENT FILING DATE: 2002-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: 22105, A NOVEL HUMAN THIOREDOXIN FAMILY TITLE OF INVENTION: MEMBER AND USES THEREOF FILE REFERENCE: 10448-022001
                                                                                                                                                                                               Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0
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OTHER INFORMATION: unsure at all Xaa locations
FEATURE:
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                           REFERENCE: 10448-188001
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Pred. No. 6.2e+0
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US-10-424-599-166951
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US-10-437-963-164055
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                                                                           APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 166951
LENGTH: 53
TYPE: PRT
ORGANISM: Glycine max
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SEQ ID NO 164055
LENGTH: 52
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Publication No. US/20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
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APPLICANT: La Rosa, Thomas
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Best Local (
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Best Local Similarity
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CURRENT FILING DATE: 2003-05-14
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                                             FEATURE: OTHER INFORMATION: Clone ID: PAT_MRT3847_121771C.1.pep
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                                                                                                                                                                                                                                                                                     Kovalic David K
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100.0%;
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Score 31;
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DB 4; Length 53;
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RESULT 48
US-09-864-761-42458
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PRIOR PILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
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Matches 4; Conserv
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                           OTHER INFORMATION: NOTHER INFORMATION: EOTHER INFORMATION:
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APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/234,687
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                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                   ENGTH:
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R APPLICATION N
R FILING DATE:
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APPLICATION NUMBER: PCT/US01/00668
FTIING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 09/608,408
FILLING DATE: 2000-06-30
APPLICATION NUMBER: US 09/774,203
FILING DATE: 2001-01-29
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Hanzel, David K.
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                                                                                                                                                                     AC011510.3
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ID IN BRAIN, SIGNAL = 2.
ID IN ADULT LIVER, SIGNAL = 1.9
ID IN HEART, SIGNAL = 2.4
ID IN PLACENTA, SIGNAL = 2.4
ID IN FETAL LIVER, SIGNAL = 2.4
ID IN FETAL LIVER, SIGNAL = 4.7
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; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.7; OTHER INFORMATION: EST HUMAN HIT: BE148597.1, EVALUE 2.00e-24; OTHER INFORMATION: SWISSPROT HIT: P03204, EVALUE 3.50e+00 US-09-864-761-42458
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                                              US-10-425-115-307743
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CURRENT APPLICATION NUMBER: US/10/318,302
CURRENT FILING DATE: 2002-12-12
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin version 3.1
SEQ ID NO 2
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Best Local S
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APPLICANT: Chae,
APPLICANT: Chae,
APPLICANT: Gho,
APPLICANT: Yang,
APPLICANT: Kwon,
                                                                                                                                    NUMBER OF SEQ ID NOS:
SEQ ID NO 307743
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                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                         Sequence 307743, Application US/10425115 Publication No. US20040214272A1
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               Query Match
 Best Local Similarity
                                                                                                                                                                TITLE OF INVENTION: Nucleic Acid Molecules TITLE OF INVENTION: Plants FILE REFERENCE: 38-21(5322)B CURRENT APPLICATION NUMBER: US/10/425,115 CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                           APPLICANT: La Rosa, Thomas
APPLICANT: Kovalic, David
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bae, Dong-Goo
APPLICANT: Hwang, Sewook
TITLE OF INVENTION: BETA-AMYLOID BINDING FACTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 55
TYPE: PRT
ORGANISM: Homo sapiens
                                                         TYPE: PRT
ORGANISM: Zea mays
FEATURE:
OTHER INFORMATION: Clone ID: MRT4577_43730C.1.pep
                                                                                                                        ENGTH: 55
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Kwon, Byung Oh
Bae, Dong-Goo
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Gho, Yong Song
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-868-577A-28
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SEQ ID NO 28
LENGTH: 55
                                                           NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 26469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 26469, Application US/11097143 Publication No. US20050208558A1 GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: DETECTION KIT, SUCH AS TITLE OF INVENTION: ARRAYS, FOR DETECTING TITLE OF INVENTION: DROSOPHILA GENES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 2004-06-14
PRIOR APPLICATION NUMBER: US 60/478,390
PRIOR FILING DATE: 2003-06-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: HEPARIN BINDING VEGER-3 LIGANDS FILE REFERENCE: 28967/39359A CURRENT APPLICATION NUMBER: US/10/868,577A
                                                                                                                                PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/161,932 PRIOR FILING DATE: 1999-10-28
                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/160,191 PRIOR FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT FILING DATE: 2005-04-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: CL000728
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APPLICANT: et al.
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PRIOR FILING DATE: 2003-09-23
                    LENGTH: 59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/157,832
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ORGANISM: DROSOPHILA
                                                                                                                                                                                                           APPLICATION NUMBER: 60/173,383
FILING DATE: 1999-12-28
APPLICATION NUMBER: 60/175,693
FILING DATE: 2000-01-12
                                                                                                                                                FILING DATE: 2000-02-24
APPLICATION NUMBER: 60/191,637
                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/164,769
FILING DATE: 1999-11-12
                                                                                                                                                                                           APPLICATION NUMBER: 60/184,831
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Pred. No.
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EXPRESSION OF 10,000 OR
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TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
RUMBER OF SEQ ID NOS: 369326
SEQ ID NO 219153
LENGTH: 56
TYPE: 7-
                                                                                                                                                                                                    APPLICANT: Venter, J. Craig
APPLICANT: et al.
APPLICANT: et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS NU
TITLE OF INVENTION: ARRAYS, FOR DETECTING EX:
TITLE OF INVENTION: DROSOPHILA GENES.
FILE REFERENCE: CL000728
CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: 60/157,832
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/160,191
PRIOR FILING DATE: 1999-10-28
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR FILING DATE: 1999-10-28
PRIOR APPLICATION NUMBER: 60/164,769
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US-11-097-143-26490
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US-10-425-115-219153
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                                                           PRIOR FILING DATE: 1999-11-12
PRIOR APPLICATION NUMBER: 60/173,383
PRIOR FILING DATE: 1999-12-28
PRIOR APPLICATION NUMBER: 60/175,693
PRIOR APPLICATION NUMBER: 60/184,831
PRIOR APPLICATION NUMBER: 60/184,831
PRIOR PILING DATE: 2000-02-24
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Best Local Similarity
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Best Local Similarity
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ORGANISM: Zea mays
FEATURE:
                    APPLICATION NUMBER: 60/191,637 FILING DATE: 2000-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          45 CGPC 48
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100.0%; Pred. No. 6.8e+02;
7ative 0; Mismatches 0;
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100.0%; Pred. No. 6.7e+02;
tive 0; Mismatches 0;
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EXPRESSION OF 10,000 OR MORE
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US-10-425-115-353222
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CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 28568
                                     APPLICANT: LA ROSA, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5)222)B
FILE REFERENCE: 38-21(5)222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 353222
LENGTH: 57
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                                                                                                                                                                                                                                                                                                 Sequence 353222, Application US/10425115 Publication No. US20040214272A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 28568, Application US/10029386
Publication No. US20030194704A1
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Best Local :
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APPLICANT: Hanzel, David K.
TITLE OF INVENTION: HUVAN GENOME-DERIVED
TITLE OF INVENTION: EXPRESSION ANALYSIS
FILE REFERENCE: ABOMICA-X-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PEATURE:

OTHER INFORMATION: MAP TO CHR2.1

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.5

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.71

OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 2.7

OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.7

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.3

OTHER INFORMATION: SWISSPROT HIT: P00203, EVALUE 1.20e+00
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ORGANISM: Zea mays
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Similarity 100.0%;
4; Conservative 0.
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Pred. No. 6.8e+02;
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US-10-424-599-185374
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                                      ; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_97667C.1.pep
US-10-437-963-202381
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                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 202381
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SEQ ID NO 185374
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Publication No. US20040031072A1
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APPLICANT: La ROSa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
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Publication No. US20040123343A1
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Best Local Similarity
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)8
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
Query Match
                                                                                                                                                                                                                     APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Fing
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLS OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
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OTHER INFORMATION: Clone ID: MRT4577_85311C.1.pep
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ORGANISM: Glycine max
FEATURE:
                                                                                        TYPE: PRT
ORGANISM: Oryza sativa
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Wu, Wei
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100.0%; Pred. No. 7e-
tive 0; Mismatches
    Score 31;
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    DB 4;
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    Length 58;
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RESULT 61
US-11-097-143-10200
; Sequence 10200, Application US/11097143
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US-09-864-408A-7494
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Best Local Similarity
Marches 4; Conserve
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; OTHER INFORMATION: Clone ID: PAT_MRT3847_130272C.1.pep
US-10-424-599-176362
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US-10-424-599-176362
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                                                                                                                                                                                                                                                                   ; ORGANISM: Homo sapiens US-09-864-408A-7494
                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 9068
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7494
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LENGTH: 59
                                                                                                                                                                                   Query Match 100.0%; Score 31; Best Local Similarity 100.0%; Pred. No. Matches 4; Conservative 0; Mismatch
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Matches 4; Conserv
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TITLE OF INVENTION: Sylucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
FULCATION NUMBER: US/10/424,599
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILLING DATE: 2003-04-28
RUMMBER OF SEQ ID NOS: 285684
                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/864,408A CURRENT FILING DATE: 2001-05-24 PRIOR APPLICATION NUMBER: 60/206,690 PRIOR FILING DATE: 2000-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: No. US20040009474A1el Human Polynucleotides and Polypeptides Enc.
FILE REFERENCE: 21402-012
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
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US-10-424-599-278438
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US-11-097-143-10200
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CURRENT APPLICATION NUMBER: US/11/097,143

CURRENT FILING DATE: 2005-04-04

PRIOR APPLICATION NUMBER: 60/157,832

PRIOR FILING DATE: 1999-10-05

PRIOR APPLICATION NUMBER: 60/160,191

PRIOR APPLICATION NUMBER: 60/161,932

PRIOR APPLICATION NUMBER: 60/161,932

PRIOR APPLICATION NUMBER: 60/164,769

PRIOR APPLICATION NUMBER: 60/164,769
                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 278438
LENGTH: 62
                                                                               Query Match
Best Local S
Matches 4
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APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
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SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 10200
LENGTH: 60
TYPE: PRT
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APPLICANT: Venter, J. Craig
APPLICANT: et al.
TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
TITLE OF INVENTION: DROSOPHILA GENES.
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 278438, Application US/10424599 Publication No. US20040031072A1
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                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: 60/191,637
PRIOR FILING DATE: 2000-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/175,693
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/184,831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1999-11-12
PRIOR APPLICATION NUMBER: 60/173,383
PRIOR FILING DATE: 1999-12-28
                                                                                                                                                                                                                                    ORGANISM: Glycine max
                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                  FEATURE:
                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 100
les 4; Conservative
28
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                                         2 CGPC 5
                                                                                    4.
  CGPC 31
                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 31; DB 6; 100.0%; Pred. No. 7.1e+02; ... Mismatches 0;
                                                                                                      100.0%; Score 31; DB 4; 100.0%; Pred. No. 7.3e+02;
                                                                                    <u>,</u>
                                                                                      Mismatches
                                                                                                                             Length 62
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                                                                                      Indels
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Query Match
Best Local Similarity
Thes 4; Conserve
                                                                    ; TYPE: PRT; ORGANISM: DROSOPHILA US-11-097-143-26472
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US-11-097-143-26472
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; OTHER INFORMATION: Clone ID: MRT4577_144792C.1.pep
US-10-425-115-233772
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                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: 60/157,832
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/160,191
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/161,932
PRIOR FILING DATE: 1999-10-28
PRIOR FILING DATE: 1999-10-28
PRIOR FILING DATE: 1999-10-28
PRIOR FILING DATE: 1999-10-28
PRIOR FILING DATE: 1999-11-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 26472, Application US/11097143
Publication No. US20050208558A1
GENERAL INFORMATION:
APPLICANT: Venter, J. Craig
APPLICANT: et al.
TITLE OF INVENTION: DRTECTION KIT, SUCH AS
TITLE OF INVENTION: ARRAYS, FOR DETECTING
TITLE OF INVENTION: DROSOPHILA GENES.
                                                                                                                                    NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 26472
LENGTH: 63
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LENGTH: 63
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Publication No. US20040214272A1
Query Match
Best Local Similarity
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules
TITLE OF INVENTION: Plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/191,637 PRIOR FILING DATE: 2000-03-23
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PRIOR FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/173,383 PRIOR FILING DATE: 1999-12-8 PRIOR APPLICATION NUMBER: 60/175,693 PRIOR FILING DATE: 2000-01-12
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ORGANISM: Zea mays
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 CGPC 5
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                                                                                                                                                                                       for Windows Version
100.0%;
Score 31; DB 6;
Pred. No. 7.4e+02;
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EXPRESSION OF 10,000 OR MORE
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                                                                                  US-10-424-599-180203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-424-599-180203
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE REPERENCE: 38-21(53222)B
FILE REPERENCE: 38-21(53222)B
FURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 336734
LENGTH: 64
LENGTH: 64
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
FEATURE:
                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53223)B CURRENT APPLICATION NUMBER: US/10/424,599 CURRENT FILING DATE: 2003-04-28 NUMBER OF SEQ ID NOS: 285684 SEQ ID NOS: 285684
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Query Match
Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: unsure
LOCATION: (1)..(64)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                             FEATURE:
OTHER INFORMATION: Clone ID:
                                                                                                                                         PEATURE:
PEATURE:
LOCATION: (1)..(65)
OTHER INFORMATION: unsure at all Xaa locations
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                                                                                                                                                                                                                                                 LENGTH: 65
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17 CGPC 20
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100.0%; Score 31; DB 4; ilarity 100.0%; Pred. No. 7.5e+0; Conservative 0; Mismatches
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                       7.5e+02;
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                                        Length 65;
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US-10-425-115-256158
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                                                                                                                                                        US-10-425-115-256158
                                                                                                                                                                                                                                                                                                                                                                                                                       Publication ....
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 256158
LENGTH: 67
TYPE: PRT
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LENGTH: 67
                                                                                              Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT FAPPLICATION NUMBER: US/10/437,963
CURRENT FAPPLICATION NUMBER: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Zhou, Yihua
APPLICANT: CZO, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
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                                                                                                                                                                                             ORGANISM: Zea mays FEATURE:
                                                                                                                                                                        OTHER INFORMATION: Clone ID: MRT4577_165207C.1.pep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Oryza sativa
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Local Similarity 100.0%;
hes 4; Conservative 0;
                                                                                                Local Similarity
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10 CGPC 13
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                                       2 CGPC 5
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Barbazuk, Brad
                                                                              Conservative
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                                                                                              100.0%; Score 31; DB 4;
100.0%; Pred. No. 7.7e+02;
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US-11-097-143-32565
                                                           FILE REFERENCE: 807A
CURRENT APPLICATION NUMBER: US/10/243,552
CURRENT FILING DATE: 2002-09-12
PRIOR APPLICATION NUMBER: US 60/322,511
PRIOR FILING DATE: 2001-09-13
PRIOR PPLICATION NUMBER: PCT/US00/35017
PRIOR FILING DATE: 2000-12-22
PRIOR PILING DATE: 2000-12-25
PRIOR APPLICATION NUMBER: US 09/488,725
PRIOR PILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: US 09/487,317
PRIOR APPLICATION NUMBER: US 09/552,317
PRIOR FILING DATE: 2000-04-25
PRIOR FILING DATE: 2000-04-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Tang, Y. Tom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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APPLICANT: Wang, Zhiwei
APPLICANT: Weng, Gezhi
APPLICANT: Weng, Gezhi
TITLE OF INVENTION: Novel Nucleic Acids and
TITLE OF INVENTION: Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: CL000728
CURRENT APPLICATION NUMBER: US/11/097,143
CURRENT FILING DATE: 2005-04-04
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TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
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PRIOR APPLICATION NUMBER: 60/164,769
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APPLICATION NUMBER: US 09/491,404
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PRIOR RILING DATE: 2000-02-03
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APPLICANT: Mite, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION. Mucolytic and Anti-Elastase Compounds and Methods of Use of the Current Application NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/469,960
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/560,118
PRIOR APPLICATION NUMBER: 60/566,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2003-02-07
NUMBER OF SEQ ID NOS: 15
SOFTMARE: Patentin version 3.3
SEQ ID NO 1
LENGTH: 4
TYPE: PRT
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                                                                                                                                                                                             ; TYPE: PRT
; ORGANISM: Artificial sequence
; PEATURE:
; OTHER INFORMATION: synthetic peptide motif
US-11-090-916-1
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US-11-090-916-1
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Best Local :
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US-11-183-136-38
US-11-072-512-2161
US-11-098-686-11331
US-10-980-388-40
US-10-980-388-40
US-10-12-289-3
US-11-022-289-8
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US-10-453-372-236
US-10-453-372-236
US-10-453-372-240
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US-11-018-868-30
US-11-103-957-43
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US-11-124-367A-314
US-11-008-727-18
US-11-077-386-25
US-11-183-136-38
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RESULT 4
US-11-223-547-1
; Sequence 1, Application US/11223547
; Publication No. US20060009527A1
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RESULT

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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Marks, Paul A.

APPLICANT: Marks, Paul A.

APPLICANT: Marks, Paul A.

APPLICANT: Butler, Lisa M.

TITLE OF INVENTION: Method of Treating TRX Mediated Dis

FILE REFERENCE: 3254.1005-001

CURRENT APPLICATION NUMBER: US/11/223,405

CURRENT FILING DATE: 2005-09-09

PRIOR APPLICATION NUMBER: US/10/369,094

PRIOR FILING DATE: 2003-02-14

PRIOR FILING DATE: 2002-02-15

NUMBER OF SEQ ID NOS: 18

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/11/144,301A
CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US 60/577,089
PRIOR FILING DATE: 2004-06-04
PRIOR PILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin version 3.3
SEQ ID NO 8
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FEATURE:
OTHER INFORMATION: Synthetic peptide
US-11-144-301A-8
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US-11-223-405-1
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                                                                              Query Match 100.0%; Score 31;
Best Local Similarity 100.0%; Pred. No.
Matches 4; Conservative 0; Mismatch
                                                                                  Matches
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APPLICANT: Johanna, Ungerstedt
TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS
FILE REFERENCE: 28517-501 UTIL
                                                                                                                                                                                                            LENGTH: 4
TYPE: PRT
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                                                                                                                                                                                         ORGANISM: homo sapiens
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hes 0;
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GENERAL INFORMATION:

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Matches 4
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Best Local Similarity
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Publication No. US20060030022A1
                                                                                                                                                                                                                                                                                  SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: RITZ, DANIEL
APPLICANT: LIM, JACKE EUN-AH
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: BOND CONTAINING PROTEINS IN
FILE REFERENCE: HMV-052.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: U.S. 60/357,383
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 18
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CURRENT FILING DATE: 2005-09-09
PRIOR APPLICATION NUMBER: US/10/369,094
                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: 60/163,939
PRIOR FILING DATE: 1999-11-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US/09/679,705
PRIOR FILING DATE: 2000-10-05
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CURRENT FILING DATE: 2005-02-16
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APPLICANT: ASLUND, FREDRIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Butler, Lisa M.
TITLE OF INVENTION: Method of Treating TRX Mediated Diseases
                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/157,770
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                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                     NUMBER OF
                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/166,044
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                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: variant motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: homo sapiens
                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                       FEATURE:
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                                                                                        Similarity
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SEQ ID NOS: 24
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Pred. No. 8.2e+04;
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                                                                 PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR PELICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR PILING DATE: 2004-03-24
PRIOR PILING DATE: 2005-02-07
SOFTWARE: I
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Publication No. US20050260140A1
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Best Local Similarity
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Publication No. US20050260140A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE:
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NAME/KEY: misc_feature
LOCATION: (1)...(6)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10
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CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
                                                                                                                                                                                                                                                                                                                                                                        RPPLICANT: Lee, Rees
FITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: synthetic peptide motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial sequence FEATURE:
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                          PatentIn version 3.3
                                                  SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PatentIn version 3.3
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                                                                                                                                                                                                                                                                                                                                                     2879-98-
                                                                                                                                                                                                                                                                                                                                                                                                Rees
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US-10-467-657-5372
; Sequence 5372, Application US/10467657
; Publication No. US20050260581A1
; GENERAL INFORMATION:
; APPLICANT: CHIRON SpA
; APPLICANT: FONTANA Maria Rita
; APPLICANT: FIZZA Mariagrazia
; APPLICANT: MASIGNANI Vega
; APPLICANT: MASIGNANI Vega
; APPLICANT: MONACI Eliasbetta
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PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR FILING DATE: 2004-10-26
PRIOR FILING DATE: 2004-10-26
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Best Local Similarity 100.0%;
Matches 4; Conservative 0
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Best Local Similarity
Matches 4; Conserv
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LENGTH: 75
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PRIOR FILING DATE: 2004-10-26
NUMBER OF SEQ ID NOS: 9
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CURRENT FILING DATE: 2005-05-17
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APPLICANT: LEITE, JOHN F.
TITLE OF INVENTION: COMPOSITIONS,
TITLE OF INVENTION: SPECTROMETRY
FILE REFERENCE: NVX-28 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: HAJIVANDI,
APPLICANT: POPE, ROBI
APPLICANT: SADEGHI, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 3.3
CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
                                         TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Artificial Sequence
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SADEGHI, MEHRNOOSH
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Pred. No.
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NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn Ver. 3.3
SEQ ID NO 1
LENGTH: 91
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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CURRENT FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR APPLICATION NUMBER: 60/621,686
PRIOR APPLICATION NUMBER: 60/621,686
PRIOR APPLICATION NUMBER: 60/621,686
PRIOR APPLICATION NUMBER: 60/621,686
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APPLICANT: HAJIVANDI, MAHBOD R.
APPLICANT: POPE, ROBERT M.
APPLICANT: SADEGHI, MEHRNOOSH
APPLICANT: LIANG, XIQUAN
APPLICANT: LIANG, YIQUAN
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PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
SOFTWARE: SeqWin99, version 1.04
SEQ ID NO 5372
LENGTH: 84
                                                                             Sequence 286, Application No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1, Application US/11131744
Publication No. US20060023808A1
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Best Local Similarity
Matches 4; Conserv
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APPLICANT: LEITE, JOHN F.

TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR CALIBRATION IN MASS

TITLE OF INVENTION: SPECTROMETRY

FILE REFERENCE: NVX-28 US
APPLICANT: Klaenhammer, Todd R. APPLICANT: Russell, William M. APPLICANT: Altermann, Eric APPLICANT: McAuliffe, Olivia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 2004-10-26
                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
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1 Similarity 100.0%;
4; Conservative 0;
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                                                                                                                       Application US/11074176
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                    US20050250135A1
                                                                                                                                                                                                                                                                                                                           100.0%; Score 31;
100.0%; Pred. No.
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Pred. No.
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RESULT 13
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Sequence 7, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
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LENGTH: 103
TYPE: PRT
ORGANISM: Lactobacillus acidophilus
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Best Local
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Best Local 9
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APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
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CURRENT FILING DATE: 2005-03-24
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SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT APPLICATION NUMBER: US/11/074,176
CURRENT FILING DATE: 2005-03-07
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PRIOR APPLICATION NUMBER: 60/556,516
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PRIOR APPLICATION NUMBER: 10/660,118
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TITLE OF INVENTION: Nucleic Acid Sequences Encoding
TITLE OF INVENTION: Stress-Related Proteins and Uses Therefore
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/650,865
                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Listeria monocytogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 103
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PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR FILING DATE: 2003-09-10
PRIOR FILING DATE: 2004-03-24
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
PRIOR FILING DATE: 2005-02-07
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Best Local S
Matches 4
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PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-04-11
PRIOR FILING DATE: 2003-09-10
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/650,865
                                                                                                                                                                                                                        SOFTWARE: PatentIn version 3.3 SEQ ID NO 5
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Publication No. US20050260140A1
GENERAL INFORMATION:
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Best Local Similarity
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TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
CURRENT FILING DATE: 2005-03-24
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APPLICANT: del Va
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CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
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TYPE: PRT
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TYPE: PRT
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                                                                                          Local Similarity
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ilarity 100.0%;
Conservative 0
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RESULT 15 US-10-621-234-1371 ; Sequence 1371, Application US/10821234 ; Publication No. US20050255114A1

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RESULT 17
US-11-090-916-9
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US-10-821-234-1371
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PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR PELICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
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CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
SOFTWARE: pt SEQ genes Version 1.0
SEQ ID NO 1371
                 Sequence 9, Application US/11090916 Publication No. US20050260140A1 GENERAL INFORMATION:
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Publication No. US20050260140A1
GENERAL INFORMATION:
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Best Local Similarity
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PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
SOPTWARE: PatentIn version 3.3
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APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
APPLICANT: White, Carl W.
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CURRENT FILING DATE: 2005-03-24
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TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
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CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2002-09-10
PRIOR PRIOR DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/550,865
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
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Matches
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                                                                                                                                                                                                                                                                                                                                                   APPLICANT: White, Carl W.
APPLICANT: White, Carl W.
APPLICANT: Lee, Rees
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use TITLE OF INVENTION: MUCOLYTIC and Anti-Elastase Compounds and Methods of Use TILE REFERENCE: 2879-98-1
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
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SEQ ID NO 9
                                                                               Matches
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                                                                                                  Best Local Similarity
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PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
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                                                                                                                                                                                LENGTH: 105
TYPE: PRT
ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Mus musculus
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100.0%; Pred. No. 32
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; ORGANISM: Homo sapiens
US-11-090-916-12
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US-11-090-916-11
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PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.3
SEQ ID NO 12
LENGTH: 105
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                                                         Query Match 100.0%;
Best Local Similarity 100.0%;
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Best Local Similarity
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Publication No. US20050260140A1
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PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
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CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 10/660,118
PRIOR TLING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
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PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
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TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
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APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
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NUMBER OF SEQ ID NOS: 15
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                                       Conservative
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Pred. No.
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US-11-144-301A-10
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US-11-144-301A-9
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US-11-106-796-13
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LENGTH: 105
TYPE: PRT
Sequence 10, Application US/11144301A Publication No. US20050288227A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn version 3.3
SEQ ID NO 9
LENGTH: 105
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APPLICANT: Marks, Paul A.
APPLICANT: Johanna, Ungerstedt
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Publication No. US20050282191A1
GENERAL INFORMATION:
APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 4; Conserv
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TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
FILE REFERENCE: 28517-501 UTIL
CURRENT APPLICATION NUMBER: US/11/144,301A
CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US 60/577,089
PRIOR FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR PILING DATE: 2003-02-14
PRIOR PILING DATE: 2003-02-15
PRIOR FILING DATE: 2002-02-15
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CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: Marks, Paul A.

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US-11-032-773-943
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Matches
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NUMBER OF SEQ ID NOS: 958
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 943
LENGTH: 105
                                                                           Sequence 19, Application US/10689742 Publication No. US20050250180A1 GENERAL INFORMATION:
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Publication No. US20
GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity
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PRIOR FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 10
APPLICANT: Jacobs, Kenneth
APPLICANT: McCoy, John M
APPLICANT: LaVallie, Edward R
APPLICANT: Racie, Lisa A
                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank BAA04881
DATABASE ENTRY DATE: 2002-12-25
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TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
FILE REFERENCE: 28517-501 UTIL
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100.0%; Pred. No. 32;
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APPLICANT: Kapur, Vivek and Gebhart, Connie J.
TITLE OF INVENTION: NUCLEIC ACID AND POLYPEPTIDE SEQUENCES
TITLE OF INVENTION: FROM LAWSONIA INTRACELLULARIS AND METHODS
FILE REFERENCE: 09531-128001
CURRENT APPLICATION NUMBER: US/11/098,686
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US03/31318
PRIOR APPLICATION NUMBER: PCT/US03/31318
PRIOR FILING DATE: 2003-10-01
PRIOR FILING DATE: 2003-10-01
PRIOR FILING DATE: 2003-10-04
NUMBER OF SEQ ID NOS: 11433
SOPTWARE: FRASESEQ for Windows Version 4.0
SEQ ID NO 10612
LENGTH: 107
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US-11-098-686-10612
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Sequence 4, Application US/11090916
Publication No. US20050280140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
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Publication No. US20060024696A1
GENERAL INFORMATION:
APPLICANT: Kapur, Vivek and Gebhart, Co
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Best Local Similarity
Matches 4; Conserv
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CURRENT FILING DATE: 2003-10-22
PRIOR APPLICATION NUMBER: 09/746,783
PRIOR FILING DATE: 2000-12-21
NUMBER OF SEQ ID NOS: 231
SOFTWARE: PatentIn version 3.2
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APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: 00766.000091.10
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TYPE: PRT
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100.0%; Pred. No. 33,
ative 0; Mismatches
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                                                                                                                            ; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-072-512-3199
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SEQ ID NO 4
LENGTH: 109
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Best Local
                                                                 Matches
                                                                               Query Match
Best Local Similarity
                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3199
LENGTH: 113
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Publication No. US20060029945A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 10/660,118
PRIOR PILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR PILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR PILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/11/072,512
CURRENT FILING DATE: 2005-03-07
EVICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2002-01-25
PRIOR FILING DATE: 2002-01-25
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APPLICANT:
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NUMBER OF SEQ ID NOS: 4096
                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: JP 2001-379298
                                                                                                                                                                                                                                                                                                                                               APPLICANT: NAGAHARI, KENJI
APPLICANT: MASUHO, YASUHIKO
TITLE OF INVENTION: Novel full length cDNA
FILE REFERENCE: 084335-0191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: ISOGAI, TAKAO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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37 CGPC 40
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                               2 CGPC 5
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NAGAI, KEIICHI
IRIE, RYOTARO
TAMECHIKA, ICHIRO
SEKI, NAOHIKO
YOSHIKAWA, TSUTOMU
OTSUKA, MOTOYUKI
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OTSUKI, TETSUJI
WAKAMATSU, AI
SATO, HIROYUKI
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YURI
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YAMAMOTO, JUN-ICHI
                                                                 Conservative
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                                                                               100.0%; Score 31; DB 100.0%; Pred. No. 34;
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                                                               Mismatches
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                                                                                               DB 7; Length 113;
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                                                                                                                    ; LENGTH: 122
; TYPE: PRT
; ORGANISMF Hordeum vulgare
US-11-195-459-16
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CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
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                                                                                                                                                                                                  SEQ ID NO 16
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Publication No. US20050278803A1
                                         Matches
                                                           Query Match
Best Local Similarity
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Best Local Similarity
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APPLICANT: Everard, John
APPLICANT: Allen, Stephen
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PROFILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Sewalt, Vincent
APPLICANT: Hastings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Sewalt, Vincent APPLICANT: Hastings, Crai
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    2 CGPC 5
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Hantke, Sabine
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                                      Conservative
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                                      100.0%; Score 31; D1
100.0%; Pred. No. 36
tive 0; Mismatches
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Pred. No. 36;
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                                                         DB 7; Length 122; 36;
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FEATURE:
FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: "n" at position 9, 493, 537, 548,
OTHER INFORMATION: or t
US-11-195-459-19
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SEQ ID NO 10
LENGTH: 123
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Query Match
Best Local Similarity
Matches 4; Conserv
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SEQ ID NO 19
LENGTH: 122
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Best Local (
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                                                                                                                                                                                                                                                         APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PROFILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILLING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILLING DATE: 2001-12-03
PRIOR FILLING DATE: 2001-12-03
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE
PILE REFERENCE: 5718-119 (035718/241421)
CURRENT EPPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-03
                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sewalt, Vincent
APPLICANT: Hastings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
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PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
                                                                                                    TYPE: PRT
ORGANISM: Zea mays
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No. US20050278803A1
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No. US20050278803A1
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    100.0%; Score 31; DB 7; Length 123; ilarity 100.0%; Pred. No. 36; Conservative 0; Mismatches 0; Indels
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TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METH
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 703
LENGTH: 126
TYPR: DOT
RESULT 35
US-11-106-796-11
; Sequence 11, Application US/11106796
; Publication No. US20050282191A1
; GENERAL INFORMATION:
; APPLICANT: SUTOVSKY, PETER
; APPLICANT: MITANDA-VIZUETE, ANTONIO
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
                                                                                                                                                                                                                                                                                                                                                            US-11-106-796-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Homo sapiens US-10-995-561-703
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US-10-995-561-703
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                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 10, Application US/11106796
Publication No. US20050282191A1
GENERAL INFORMATION:
APPLICANT: SUTOYSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 703, Application US/10995561
Publication No. US20050272054A1
GENERAL INFORWATION:
APPLICANT: CARGILL, Michele et al.
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Best Local (
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CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 127
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100.0%; Pred. No. 37
tive 0; Mismatches
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; TYPE: PRT
; ORGANISM: Mus musculus
US-11-106-796-11
              SOFTWARE: SeqWin99,
SEQ ID NO 4354
LENGTH: 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 12
LENGTH: 127
TYPE: PRT
ORGANISM: RAT
                                                                                                                                                                                                                                                                           Sequence 4354, Application US/10467657 Publication No. US20050260581A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 4; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 12, Application No. US20 GENERAL INFORMATION:
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Best Local Similarity
                                                                  CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
PRIOR APPLICATION NUMBER: GB-0103424.8
PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
                                                                                                                                                                      APPLICANT: CHIRON SPA
APPLICANT: FONTANA MARIA Rita
APPLICANT: FIZZA MARIAGRAZIA
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Elisabetta
APPLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
FILE REFERENCE: UNMO:033US
CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
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US-11-072-512-2145
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Matches
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PRIOR APPLICATION NUMBER: 60/409,960
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                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
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APPLICANT: MASUHO, YASUHIKO
TITLE OF INVENTION: Novel full length cDNA
FILE REFERENCE: 084335-0191
CURRENT APPLICATION UNMERR: US/11/072,512
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION UNMERR: US 60/350,978
PRIOR FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: JP 2001-379298
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 4096
                                                                                                                               Sequence 13, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2145
LENGTH: 129
TYPE: PRT
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Best Local Similarity 100.
Matches 4; Conservative
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APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: del Res
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Neisseria gonorrhoeae
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NAGAI, KEIICHI
IRIE, RYOTARO
TAMECHIKA, ICHIRO
SEKI, NAOHIKO
YOSHIKAWA, TSUTOMU
OTSUKA, MOTOYUKI
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OTSUKI, TETSUJI
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YAMAMOTO, JUN-ICHI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WAKAMATSU, AI
SATO, HIROYUKI
                                                                                                                                                                                                                                                                                                                                             100.0%; Score 31; Dilarity 100.0%; Pred. No. 37, Conservative 0; Mismatches
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Pred. No.
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37;
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Query Match
Best Local Similarity
Watches 4; Conserva
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US-11-195-459-8
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                                                                                                                       RESULT 41
US-09-978-360A-641
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                                 Sequence 641, Application US/09978360A
Publication No. US20060009633A9
GENERAL INFORMATION:
APPLICANT: Edwards, Jean-Baptiste Dumas Milne
APPLICANT: Duclert, Aymeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Sewalt,
APPLICANT: Hasting
APPLICANT: Meeley,
APPLICANT: Hantke,
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Best Local
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APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/95,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-03
                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
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   APPLICANT:
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TYPE: PRT
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APPLICATION NUMBER: 60/556,516
FILING DATE: 2004-03-24
APPLICATION NUMBER: 60/650,865
FILING DATE: 2005-02-07
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Meeley, Robert
Hantke, Sabine
Duclert, Aymeric
Bougueleret, Lydie
Jobert, Severin
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Pred. No. 38;
0; Mismatches
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Pred. No.
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APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: del Val, Greg
APPLICANT: del Val, Greg
ITILE DF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 10/60,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
CONTROL OF SEQ ID NOS: 15
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; LOCATION: -41..-1
US-09-978-360A-641
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PRIOR FILING DATE: 1997-12-17
PRIOR PELICATION NUMBER: US 60/074,121
PRIOR FILING DATE: 1998-02-09
PRIOR APPLICATION NUMBER: US 60/081,563
PRIOR FILING DATE: 1998-04-13
PRIOR PELICATION NUMBER: US 60/096,116
PRIOR FILING DATE: 1998-08-10
PRIOR FILING DATE: 1998-08-10
PRIOR FILING DATE: 1998-08-10
PRIOR APPLICATION NUMBER: US 60/099,273
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 810

SOFTWARE: Patent.pm

SEQ ID NO 641
LENGTH: 143
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Matches
                   SOFTWARE: PatentIn version 3.3 SEQ ID NO 14 LENGTH: 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14, Application US/11090916 Publication No. US20050260140A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: -09-04
PRIOR APPLICATION NUMBER: US 09/191,997
PRIOR FILING DATE: 1998-11-13
PRIOR APPLICATION NUMBER: US 09/215,435
PRIOR FILING DATE: 1998-12-17
PRIOR FILING DATE: 1998-12-17
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CURRENT FILING DATE: 2001-10-15
PRIOR APPLICATION NUMBER: US 60/66,677
PRIOR FILING DATE: 1997-11-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
TYPE: PRT
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FILING DATE: 1998-12-17
APPLICATION NUMBER: US 09/247,155
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ORGANISM: Zea mays

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APPLICANT: LIANG, XIQUAN
APPLICANT: LEITE, JOHN F.
ITITLE OF INVENTION: COMPOSITIONS, KITS, AND
ITITLE OF INVENTION: SPECTROMETRY
FILE REFERENCE: NVX-28 US
CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/69,373
PRIOR APPLICATION NUMBER: 60/69,373
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
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US-11-090-916-15
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PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2003-09-10
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Publication No. US20060023808A1
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Best Local Similarity
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APPLICANT:
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
                                                                                                                                                                                                                                                                                                  APPLICANT: HAJIVANDI, MAHBOD R. APPLICANT: POPE, ROBERT M.
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PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Oryza sativa
FILING DATE: 2004-05-17
APPLICATION NUMBER: 60/621,685
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                                                                                                                                                                                                                     METHODS FOR CALIBRATION IN MASS
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TITLE OF INVENTION: GENETIC POLYMORPHISMS AS TITLE OF INVENTION: CARDIOVASCULAR DISORDER TITLE OF INVENTION: DETECTION AND USES THER FILE REFERENCE: CL001559
FULE REFERENCE: CL001559
FULE REPERENCE: CL001559
FULE REFERENCE: CL001559
FULL REFERENC
                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 46
US-10-995-561-643
US-10-995-561-643; Sequence 643, Application US/10995561; Publication No. US20050272054A1; PUBLICANT: CARGILL, Michele et al.; APPLICANT: CARGILL, Michele et al.; APPLICANT: CARGILL, Michele et al.
; TYPE: PRT; ORGANISM: Homo sapiens US-10-995-561-643
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SEQ ID NO 3
LENGTH: 177
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
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Best Local Similarity
Matches 4; Conserv
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SOFTWARE: PatentIn Vers. 2.0
SEQ ID NO 6
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APPLICANT: Pleach,
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Best Local Similarity 100.0%; Pred. No. 45
Matches 4; Conservative 0; Mismatches
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CURRENT APPLICATION NUMBER: US/10/467,962B
CURRENT FILING DATE: 2003-08-14
PRIOR APPLICATION NUMBER: PCT/EP02/01466
PRIOR FILING DATE: 2002-02-13
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APPLICANT: Daeschner, Klaus
APPLICANT: Klain, Mathieu
TITLE OF INVENTION: Identification of Herbicidally Active Substances
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ORGANISM: Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH: 183
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                                                                                                                                                                                                                                                                                                                                                                                Michele et al.
GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RES
DETECTION AND USES THEREOF
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Pred. No.
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RESULT 47
US-10-995-561-642
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; ORGANISM: Homo sapiens
US-10-995-561-642
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GENERAL INFORMATION:
APPLICANT: CARGILL, Michele et al.
APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CL001.559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT APPLICATION NUMBER: US/10-295,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 642
LENGTH: 191
                                                                                                                                                                                                                                                                      US-11-064-774A-2
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Best Local S
Matches 4
                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: MATERIALS AND METHODS INVOLVING HYBRID VASCULAR ENDOTHELIAL GROWT
TITLE OF INVENTION: PACTOR DAS AND PROTEINS
FILE REFERENCE: 28967/35977B2
CURRENT APPLICATION NUMBER: US/11/064,774A
CURRENT FILING DATE: 2005-02-24
PRIOR APPLICATION NUMBER: 09/795,006
PRIOR FILING DATE: 2001-02-26
PRIOR APPLICATION NUMBER: US 60/205,331
PRIOR APPLICATION NUMBER: US 60/205,331
PRIOR FILING DATE: 2000-05-18
PRIOR FILING D
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Best Local Similarity
Matches 4; Conserv
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Best Local Similarity
                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Alitalo et al.
                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 191
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   143 CGPC 146
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                                                                                                                               Conservative
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                                                                                                                                                              100.0%;
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                                                                                                                                                                  Score 31; DB
Pred. No. 47;
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                                                                                                                                                                                              DB 7; Length 191;
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                                                                                                                                   0;
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; SEQ ID NO 2
; LENGTH: 191
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-075-400-2
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US-11-226-005-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4, Application US/11226005 Publication No. US20060014252A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: ERIKKSON, et al.
TITLE OF INVENTION: MULTIVALENT ANTIBODY MATERIALS AND METHODS FORVEGF/PDGF FAMILY OF
TITLE OF INVENTION: GROWTH FACTORS
FILE REFERENCE: 28967/39820B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 32
                TELEX: 756822
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                   COMPUTER: Apple Macintosh
ODERATING SYSTEM: Apple 7.1
SOFTWARE: Microsoft Word, Version 5.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/226,005
FILING DATE: 14-Sep-2005
CIASSIFICATION DATA:
APPLICATION UNMBER: US/08/743,868
APPLICATION UNMBER: US/08/743,868
FILING DATE: 05-NOV-1996
APPLICATION UNMBER: 08/54,374
FILING DATE: 08-NOV-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                              REFERENCE/DOCKET NUMBER: 2837
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lyman, Stewart D. et al.
TITLE OF INVENTION: flk-1 Binding Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Immunex Corporation STREET: 51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Seattle
                                                                                                                                                                               NAME: Anderson, Kathryn A. REGISTRATION NUMBER: 32,172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: US
                                                                                        TELEPHONE: (206) 587-04
TELEFAX: (206) 233-0644
ENGTH: 191 amino acids
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; SEQ ID NO 9
; LENGTH: 191
; TYPE: PRT
; ORCANISM: Homo sapiens
US-11-129-076-9
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당
; ORGANISM: Bovine US-11-145-494-2
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US-11-129-076-9
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Best Local Similarity lov.
"---hes 4; Conservative
                                SEQ ID NO 2
LENGTH: 191
TYPE: PRT
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Publication No. US20060019891A1
                                                                                                                                                                                                                                                                                                                                                                  Sequence 2, Application US/11145494 Publication No. US20060024785A1 GENERAL INFORMATION:
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APPLICANT: Edelberg, Jay A.
APPLICANT: Xaymarden, Munirva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
                                                                                                                  PRIOR APPLICATION NUMBER: US/09/346,069
PRIOR FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: 08/567,200
PRIOR EILING DATE: 1995-12-05
PRIOR APPLICATION NUMBER: 60/002,827
PRIOR FILING DATE: 1995-08-25
                                                                                                                                                                                                                      FILE REFERENCE: A-62326-2
CURRENT APPLICATION NUMBER: US/11/145,494
CURRENT FILING DATE: 2005-06-03
                                                                                                                                                                                                                                                                           APPLICANT: Keyt, Bruce A.
APPLICANT: Nguyen, Francis H.
APPLICANT: Ferrara, Napoleone
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING VARIANTS OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: CELL GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/11/129,076
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Xaymarden, Munirva
APPLICANT: Cornell Research Foundation, I
TITLE OF INVENTION: Protection of Cardiac
FILE REFERENCE: 1676.009US1
                                                                                     NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IMMEDIATE SOURCE:
CLONE: Hu VEGF
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein HYPOTHETICAL: NO
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Pred. No. 47;
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PRIOR APPLICATION NUMBER: US 60/416,395
PRIOR FILING DATE: 2002-10-04
NUMBER OF SEQ ID NOS: 11433
SOPTWARE: FRACTSC for Windows Version 4.0
SEQ ID NO 11017
LENGTH: 197
TYPE: PRT
ORGANISM: Lawsonia intracellularis
US-11-098-686-11017
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US-11-098-686-11017
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; ORGANISM: Homo sapiens
US-11-075-047A-81
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                                                                                                                                                                                                                                                                            Sequence 11017, Application US/11098686

Publication No. US20060024696A1

GENERAL INFORMATION:

APPLICANT: Kapur, Vivek and Gebhart, Connie J.

APPLICANT: NUCLEIC ACID AND POLYPEPTIDE SEQUENCES

TITLE OF INVENTION: FROM LAWSONIA INTRACELLULARIS AND METHODS OF USING

FILE REFERENCE: 09531-128001

CURRENT APPLICATION NUMBER: US/11/098,686

CURRENT FILING DATE: 2003-04-04

PRIOR APPLICATION NUMBER: PCT/US03/31318

PRIOR FILING DATE: 2003-10-01
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SOFTWARE: Patentin version 3.2
SEQ ID NO 81
LENGTH: 191
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Publication No. US20060030000A1
GENERAL INFORMATION:
APPLICANT: ALITALO, et al.
TITLE OF INVENTION: GROWTH FACTOR BINDING CONSTRUCTS MATERIALS AND METHODS
FILE REFERENCE: 28967/39700A
CURRENT APPLICATION NUMBER: US/11/075,047A
CURRENT FILING DATE: 2004-03-07
PRIOR APPLICATION NUMBER: US 60/550,907
PRIOR FILING DATE: 2004-03-07
                                                                Matches
                                                                                               Query Match
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Best Local Similarity 100.0%;
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                                                                               Local Similarity
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                                                                               Score 31;
Pred. No.
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                                                                                               DB 7;
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RESULT 55 US-10-467-657-8494 ; Sequence 8494, Application US/10467657

Publication No. US20050260581A1

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GENERAL INFORMATION:

APPLICANT: CARGILL, Michele et al.

TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH

TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF

TITLE OF INVENTION: DETECTION AND USES THEREOF

FILE REFERENCE: CL001559

CUURRENT APPLICATION NUMBER: US/10/995,561

CUURRENT FILING DATE: 2004-11-24

NUMBER OF SEQ ID NOS: 85702

SOUTWARE: FRSHSEQ for Windows Version 4.0

SEQ ID NO 640

LENGTH: 209

TYPE: PRT

ORGANISM: Homo sapiens
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CURRENT FILING DATE: 2003-08-11
PRIOR APPLICATION NUMBER: GB-0103424.8
PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
SOFTWARE: SeqWin99, version 1.04
SEQ ID NO 8494
LENGTH: 209
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Sequence 645, Application US/10995561
Publication No. US20050272054A1
GENERAL INFORMATION:
APPLICANT: CARCILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDICONASCULAR DISORDERS AND DRUG RESPONSE, METHODS
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FASESEQ for Windows Version 4.0
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Best Local
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Best Local Similarity
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APPLICANT: FONTANA MARIA RITA
APPLICANT: FIZZA MARIAGRAZIA
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Elisabetta
TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE:
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nes 4; Conserv
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Query Match
Best Local Similarity ...
Approximately ...
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US-10-995-561-646
                                                                                         US-11-149-462-3
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TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CRADIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 646
LENGTH: 215
                                                                                                                                                                                                                                                                                                                   APPLICANT: Shea, Lonnie D.
APPLICANT: Shea, Lonnie L.
APPLICANT: Shea, Lonnie L.
APPLICANT: Whittlesey, Kevin
APPLICANT: Yang, Yang
APPLICANT: Rives, Christopher
APPLICANT: Rovedo, Mark
APPLICANT: Iskandar, Bermans
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Best Local Similarity 1vv.
"""hes 4; Conservative
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LENGTH: 215
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LENGTH: 215
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Best Local Similarity
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                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/578,785
PRIOR FILING DATE: 2004-06-10
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/11/149,462
CURRENT FILING DATE: 2005-06-09
                                                                                                                                                                                                                                                                                 APPLICANT: Iskandar, Bermans
TITLE OF INVENTION: Biodegradable Scaffolds and Uses Thereof
FILE REFERENCE: 1720-1-011N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
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            100.0%; Score 31; DB ilarity 100.0%; Pred. No. 51; Conservative 0; Mismatches
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US-11-106-399-8
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; ORGANISM: Homo sapiens US-11-072-512-3343
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                                                                                                                                               APPLICANT: NACAHARI, KENJI
APPLICANT: NACHARI, KENJI
APPLICANT: MASUHO, YASUHIKO
TITLE OF INVENTION: NOVE1 full length cDNA
FILE REFERENCE: 084335-0191
CURRENT APPLICATION UMBER: US/11/072,512
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2002-01-25
PRIOR FILING DATE: 2002-01-25
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CURRENT APPLICATION NUMBER: US/11/106,399
CURRENT FILING DATE: 2005-04-14
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PATENTIN VEY. 2.1
SEQ ID NO 8
LENGTH: 216
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Publication No. US20060002892A1
GENERAL INFORMATION:
APPLICANT: MATHEW, PORUNELLOOR A.
APPLICANT: MOLES, KENT S.
TITLE OF INVENTION: LLT USES THEREOF IN IMMUNE SYSTEM MODULATION
                               PRIOR APPLICATION NUMBER: JP 2001-379298
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 4096
SOPTWARE: PatentIn Ver. 2.1
SEQ ID NO 3343
LENCTH: 224
TYPE: PRT
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Best Local Similarity
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APPLICANT:
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NAGAI, KEIICHI
IRIE, RYOTARO
TAMECHIKA, ICHIRO
SEKI, NAOHIKO
YOSHIKAWA, TSUTOMU
OTSUKA, MOTOYUKI
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SATO, HIROYUKI
ISHII, SHIZUKO
YAMAMOTO, JUN-ICHI
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OTSUKI, TETSUJI
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RESULT 64 US-11-153-880-7

Sequence 7, Application US/11153880 Publication No. US20050256050A1 GENERAL INFORMATION:

APPLICANT:

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APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS AS
TITLE OF INVENTION: CARDIOVASCULAR DISORDER
TITLE OF INVENTION: DETECTION AND USES THER
FILE REFERENCE: CL001559
CURRENT APPLICATION UNMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 641
LENGTH: 232
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; ORGANISM: Homo sapiens
US-10-995-561-641
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US-10-995-561-644
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Best Local Similarity 100.
                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 644
LENGTH: 232
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Publication No. US20
GENERAL INFORMATION:
                                                               Matches
                                                                              Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                            TITLE OF INVENTION: GENEY TITLE OF INVENTION: CARITITLE OF INVENTION: DETIFILE REFERENCE: CL001559
                                                                                                                                                             LENGTH: 23
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                              APPLICANT: CARGILL,
                                                                                                                                             ORGANISM: Homo sapiens
184 CGPC 187
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                              2 CGPC 5
                                                           ch 100.0%; Score 31; DB Similarity 100.0%; Pred. No. 53; 4; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                            Michele et al.
GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Michele et al.
GENETIC POLYMORPHISMS ASSOCIATED WITH
CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DETECTION AND USES THEREOF
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                                                                                             DB 6; Length 232;
                                                                                                                                                                                                                                                                                                              AND DRUG RESPONSE, METHODS
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RESULT 65
US-11-064-774A-147
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                                                                                                                                                                                                                                         Publication No. US20050267
GENERAL INFORMATION:
APPLICANT: Alitalo et al.
                                                                                                                                                                                                                                                                                                  Sequence 147, Application US/11064774A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
              CURRENT APPLICATION NUMBER: US/11/064,774A CURRENT FILING DATE: 2005-02-24 PRIOR APPLICATION NUMBER: 09/795,006 PRIOR FILING DATE: 2001-02-26 PRIOR APPLICATION NUMBER: US 60/205,331 PRIOR FILING DATE: 2000-05-18 PRIOR APPLICATION NUMBER: US 60/185,205 PRIOR APPLICATION NUMBER: US 60/185,205 PRIOR APPLICATION NUMBER: US 60/185,205 PRIOR FILING DATE: 2000-02-25
                                                                                                                                                                              TITLE OF INVENTION: MATERIALS AND METHODS INVOLVING HYBRID VASCULAR ENDOTHELIAL GROWTITLE OF INVENTION: FACTOR DNAS AND PROTEINS FILE REFERENCE: 28967/35977B2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (202)371-2540
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 232 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: CAO, LIANG
TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR 2
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08
FILING DATE: 06-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: MARKOWICZ, KAREN R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 36,351
REFERENCE/DOCKET NUMBER: 1488.1000004
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
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APPLICATION NUMBER:
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APPLICATION NUMBER:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single TOPOLOGY: Not Relevant
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SEQ ID NOS: 1212
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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; TYPE: PRT ; ORGANISM: Homo sapiens US-11-064-774A-147
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US-11-233-119-7
                                                                                                                  CURRENT FILING DATE: 205-09-23
PRIOR APPLICATION NUMBER: 09/107,997
PRIOR PELING DATE: 1998-06-30
PRIOR PELING DATE: 1998-03-13
PRIOR PELICATION NUMBER: 09/042,105
PRIOR PELICATION NUMBER: 09/999,811
PRIOR APPLICATION NUMBER: 09/999,811
PRIOR APPLICATION NUMBER: 09/99,811
PRIOR PILING DATE: 1997-12-24
PRIOR APPLICATION NUMBER: 09/207,550
PRIOR APPLICATION NUMBER: 09/207,550
PRIOR APPLICATION NUMBER: 09/207,550
PRIOR APPLICATION NUMBER: 09/207,550
PRIOR PILING DATE: 1994-03-08
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                     SEQ ID NO 7
LENGTH: 232
TYPE: PRT
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SEQ ID NO 147
LENGTH: 232
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LENGTH: 232
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Publication No. US20050287143A1
GENERAL INFORMATION:
APPLICANT: ROSEN et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7, Application US/11233119 Publication No. US20060025331A1
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US/08/465,968
PRIOR FILING DATE: 1995-06-06
PRIOR APPLICATION NUMBER: 08/207,550
PRIOR FILING DATE: 1994-03-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                              APPLICANT: HU, JING-SHAN et al,
TITLE OF INVENTION: VASCULAR ENDOTHELIAL GROWTH FACTOR
FILE REFERENCE: PF112P4D1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: PF112P1
CURRENT APPLICATION NUMBER: US/11/211,724
CURRENT FILING DATE: 2005-08-26
                                                                                SOFTWARE: PatentIn version 3.0
                                                                                                       NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Homo sapiens
ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ITLE OF INVENTION: Vascular Endothelial Growth Factor 2
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Pred. No. 53;
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US-11-233-119-7

Conservative

0

Mismatches

<u>,</u>

Indels

0

Length 232;

100.0%; Score 31; DB 7; 100.0%; Pred. No. 53;

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; NUMBER OF SEQ ID NOS: 128
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 91
; LENGTH: 232
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-075-047A-91
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US-11-131-744-6
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Best Local Similarity
Watches 4; Conserva
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Best Local Similarity
Matches 4; Conserv
                                                                               SEQ ID NO 6
LENGTH: 273
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 6, Application US/11131744
Publication No. US20060023808A1
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TITLE OF INVENTION: GROWTH FACTOR BINDING CONSTRUCTS MATERIALS AND METHODS
FILE REFERENCE: 28967/39700A
CURRENT APPLICATION NUMBER: US/11/075,047A
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US 60/550,907
PRIOR FILING DATE: 2004-03-07
                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: POPE, ROBERT M.
APPLICANT: SADECHI, MEHRNOOSH
APPLICANT: LIANG, XIQUAN
APPLICANT: LEITE, JOHN F.
TITLE OF INVENTION: COMPOSITIONS,
TITLE OF INVENTION: SPECTROMETRY
                                                                                                                                                          PRIOR APPLICATION NUMBER: 50/6 PRIOR FILING DATE: 2004-10-26
                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
                                                                                                                        SOFTWARE: PatentIn Ver. 3.3
                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/621,685
PRIOR FILING DATE: 2004-10-26
PRIOR APPLICATION NUMBER: 60/621,686
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                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2005-04-08
                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: NVX-28 US
                     ORGANISM: Artificial Sequence FEATURE:
                                                            TYPE: PRT
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
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US-11-186-284-85
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PRIOR FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US 60/339,971
PRIOR FILING DATE: 2001-12-10
PRIOR APPLICATION NUMBER: US 60/361,978
PRIOR APPLICATION NUMBER: US 60/361,978
PRIOR APPLICATION NUMBER: US 60/361,988
PRIOR APPLICATION NUMBER: US 60/381,988
PRIOR FILING DATE: 2002-05-20
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LENGTH: 282
TYPE: PRT
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Publication No.
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APPLICANT: Burgart, Lawrence J.
APPLICANT: Burgart, Lawrence J.
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND
TITLE OF INVENTION: METHODS FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND
TITLE OF INVENTION: THERAPY OF COLON CANCER
FILE REFERENCE: MPMO1-029PZNM
CURRENT APPLICATION NUMBER: US/11/186,284
CURRENT FILING DATE: 205-07-21
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Kamatkar, Shubhangi
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Monahan, John E.
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                                                                                                                                                                                                             Millis CD;
                                                                                                                                     thioredoxin analogue.
                                                                                                                                                                                                             (revised)
                                                                                                                                               (first entry)
                                  86US-00828112.
86US-00839857.
86US-00921287.
                                                                                                                                                                                                                                           cosmetics; antibiotic.
                                                      87EP-00301150
                                                                                        /note= "optional, and may optionally be joined to terminal blocking gp."
                                                                                /note= "see above"
                                                                                                     location/Qualifiers
                                                                                                                                                                                                             ADB73331
ADB8561
ADB8561
ADB856167
ADB878167
ADB87233
ADB848155
ADB35308
ADB43930
ADC49944
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Adc77895
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RESULT 2
ABP54936
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Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             It is a redox active peptide which can be used in cpds. to treat stress and ischaemia, for inhibiting lipid peroxidin. due to drugs or toxic effects, as anti-inflammatory agents, to prevent DNA damage, and to maintain the biological activity of antibiotics or proteins. Thioredoxin can be used as a general antioxidant in the food and commetic industry. It may be used in prodm. processes for antibiotics, proteins or polypeptides. Therapeutic dose is 1 microgram/kg - 100mg/kg. (Updated on 13-0CT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct to the field.)
      glutathionylacetyl)amino)phenylarsenoxide (GSAO) to peptide and protein dithiols. The results indicated that GSAO selectively binds proteins containing closely spaced thiols. The invention provides a method for the treatment and/or prophylaxis of arthritis using a compound of formula A-(L-Y)p, where A is a cell-membrane impermeable pendant group, L is a linker and/or spacer, Y is an arsenoxide or arsenoxide equivalent, and p is an integer from 1 to 10. The compound, e.g. GSAO, is useful for the treatment of arthritic conditions such as calcific periarthritis, enteropathic, chronic, thumb, hip, peripheral, psoriatic, rheumatoid, senteropathic, chronic, thumb, hip, peripheral, psoriatic, rheumatoid, senteropathic, chronic, thumb, hip, peripheral, psoriatic, rheumatoid,
                                                                                                                                                                                                                                                            Use of an arthritis
                                                                                                                                                          The present sequence is that of a hexapeptide comprising the active site sequence of thioredoxin. The hexapeptide was used in an example from the invention that examined the binding of 4-(N-(S-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thioredoxin; osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 6 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            19-MAR-2002; 2002WO-AU000310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Active site peptide of thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABP54936;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                metal catalysed oxidative damage in biological reactions and disease
                                                                                                                                                                                                                            Example 2(b); Page 42; 91pp; English.
                                                                                                                                                                                                                                                                                                             WPI; 2002-750519/81.
                                                                                                                                                                                                                                                                                                                                                                                                            19-MAR-2001; 2001AU-00003798
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                                                                                                                                                                                                                                                                                                                                           PJ,
or Jaccoud's arthritis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page
                                                                                                                                                                                                                                                            impermeable cell membrane compound
in vertebrates.
                                                                                                                                                                                                                                                                                                                                           Donoghue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organoarsenical; arsenoxide; antiarthritic; antigout; antirheumatic; arthritis; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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Pred. No.
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 gout,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
 hand or knee osteoarthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2e+06;
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RESULT 3
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Matches 6
                                                                               The present invention relates to a method for producing an antiangiogenesis plasmin fragment. The method comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin reductase, where a reduced plasmin protein is produced and the antiangiogenesis plasmin fragment having anti-angiogenesis activity, is released from the reduced plasmin protein. The plasminogen activator is preferably a urokinase-type plasminogen activator. The angiogenesis: preferably a urokinase-type plasmin reductase is selected from annexin II plasmin fragment is A61. The plasmin reductase is selected from annexin II plasmin fragment is necessary annexin II plasmin fragment is associated with a cell membrane. The method of the invention is useful for modulating (e.g. promoting or inhibiting) angiogenesis. The present sequence represents a mammalian cleavage peptide found in thioredoxin and protein disulphide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anti-angiogenesis plasmin fragment; plasminogen; plasminogen activato plasmin reductase; reduced plasmin protein; anti-angiogenesis activit urokinase-type plasminogen activator; A61; annexin II heterotetramer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    juvenile osteoarthritis, oligoarthritis, and polyarthritis (all claimed). The compound blocks angiogenesis in the synovial tissue and leucocyte ingress that triggers inflammation, with no signs or symptoms of toxicity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Producing an anti-angiogenesis plasmin fragment, useful for modulating, e.g. promoting or inhibiting angiogenesis, comprises contacting a plasminogen polypeptide with a plasminogen activator and a plasmin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Waisman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-NOV-2002; 2002US-00304287.
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protein disulphide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WAIS/) WAISMAN D. (KWON/) KWON M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2003-596985/56.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           peptide found in thioredoxin and protein disulphide isomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kwon M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in II p11 subunit; thioredoxin;
modulating angiogenesis; mamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 5;
2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plasminogen activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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Sequence 6

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RESULT 4
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ADM32929
AC ADM3
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                     The specification describes a method for increasing the liquefaction of commucus or sputum. The method comprises contacting the mucus or sputum. The method comprises contacting the mucus or sputum of the patient with a composition comprising a protein or sputum of contacted is located in the respiratory tract, gastrointestinal tract or contacted is located in the respiratory tract, gastrointestinal tract or contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total colling of sample of mucus or sputum from the patient shows a composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin. The composition further comprises nicotinamide-adenine thioredoxin. The composition further comprises nicotinamide-adenine dinucleotide phosphate (reduced form) (NADPH) for reducing the cohesiveness of, and increasing the liquefaction of excessively or cohesiveness of, and increasing the liquefaction of excessively or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Increasing the liquefaction of excessively viscous or cohesive mucus sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site
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11-APR-2003;
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2003US-0462082P
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                              Query Match
                                                                                                     treating a proliferative condition (especially colon or renal cancer) or a condition mediated by thioredoxin/thioredoxin reductase in the human or animal body. (I) are also useful for inhibiting thioredoxin/thioredoxin reductase and cell cycle progression, to regulate/inhibit cell proliferation and/or to promote apoptosis in a cell in vitro or in vivo. The present sequence represents a thioredoxin family conserved catalytic site peptide, which is given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2; 5-dienone; cytostatic; thioredoxin reductase inhibitor; proliferative condition; colon cancer; renal cancer; thioredoxin; thioredoxin reductase; cell cycle progression; cell proliferation; apoptosis; catalytic site peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. The present sequence represents an active site of a thioredoxin which is used in the method of the invention.
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                                                                                                                                                                                                               cyclohexa-2,5-dienone derivatives (I) and their salts, esters, amides, solvates, hydrates or protected forms. (I) have cytostatic activities, and can be used as thioredoxin reductase inhibitors. (I) can be used for
                                                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                      New 4-(1(sulfonyl)-1H-indol-2-yl)-4(hydroxy)-cyclohexa-2,5-dienone derivatives are thioredoxin reductase inhibitors useful for treating e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                  Stevens MFG, Westwell
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                                                                                                                                                                                                                                                               The present invention described 4-(1(Sulfonyl)-1H-indol-2-yl)-4(hydroxy)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-DEC-2002; 2002WO-GB005842.
                                                                                                                                                                                                                                                                                                                                                                                                                                                (CANC-) CANCER RES TECHNOLOGY LTD.
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l Similarity
6; Conserva
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                                                                                                                                                                                                                                                                                           Page 82; 141pp; English.
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         family conserved catalytic site peptide.
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100.0%; Pred. No.
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                                                                                                                                   The present invention describes a composition (I) which modulates the activity of a pl1 protein and effects a change in the level of cloud cell line, which involves isolating a cell. Making (M1) a CC cloud cell line, which involves isolating a cell. Then characterising the activity of a protein produced by the cell or cloud progeny of the CC cloud cell, where the protein is involved in plasminogen activation; and (2) a CC cloud cell line (II) useful in the identification of composition that CC (I) has Cytostatic activity, and can be used in antisense therapy. (I) is useful for modulating the activity of pl1 which involves administering (I) to a cell. The cell can be a human cancer cell, chosen from a HT1080 cf ibrosarcoma cell. (I) is also useful for reducing the development of cancer cell in the patient. (I) is useful for reducing the development of cancer cell in the patient. (I) is useful for inhibiting the composition that modulates pl1 activity which involves administering (I) to a cancer cell in the patient. (I) is useful for inhibiting the growth of cell cadministering (I) to a cancer cell in the patient. (I) is useful for inhibiting the growth of cell cell of a cloud cell cell in the patient. (II) is useful for identifying the composition that modulates pl1 activity which involves administering the composition that produces a change in pl1 activity is a change in the level of plasminogen activation composition that produces a change in pl1 activity. The change in the level of plasminogen activation composition that produces a change in the reamolification composition in that produces a change in pl1 activity. The composition that produces a change in pl1 activation and protein composition that produces a change in pl1 activation and protein composition that produces a change in pl1 activation accommon recommon expresents a thoredoxin and protein composition that produces a change in the examplification composition that produces a change in pl1 activation and protein composition that produces a change in
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Sequence 6
                                                                                                                       disulfide isomerase common sequence, which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; SEQ ID NO 4; 140pp; English
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Query Match

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                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a composition which modulates the activity of a pll protein and effects a change in the level of plasminogen activation by a cell. The composition is useful for reducing pll protein activity in cancer cells. They are also useful for inhibiting tumour growth or metastasis or treating cancer. The present sequence represents the amino acid sequence of the thioredoxin and protein disulphide isomerase common sequence involved with cleavage, formation and reshuffling of disulphide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New composition which modulates the activity of a p11 protein and effects a change in the level of plasminogen activation by a cell useful for treating cancer or reducing p11 protein activity in cancer cells.
                                      27-APR-1998
                                                                                                                                                                                                                                                                                                                                       Sequence 6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    p11 protein;
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Immunomodulatory peptide D7233.
                                                                        AAW45003;
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26-NOV-2002; 2002US-00304287
13-DEC-2002; 2002US-0433140P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thioredoxin and protein disulphide isomerase common sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADQ97008;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADQ97008 standard; peptide; 6 AA
                                                                                                       AAW45003 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Waisman DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-DEC-2003; 2003US-00735577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WAIS/) WAISMAN D
                                                                                                                                                                                                                                                                  Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; SEQ ID NO 4; 66pp; English.
                                                                                                                                                                                                                            WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WCGPCK 6
                                                                                                                                                                                                WCGPCK 6
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                                                                                                                                                                                                                                                                    Conservative
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                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    plasminogen
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                                                                                                       peptide; 12 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; ٢1
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                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    activation; tumour growth; metastasis;
                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. 2e+06;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                 Score 47;
Pred. No.
                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                 2e+06;
                                                                                                                                                                                                                                                                                                    DB 8;
                                                                                                                                                                                                                                                                                                  Length 6;
                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                                  0
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                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treating.
                                                                                                                                                                                                                                                                  <u>,</u>
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                                                                                                                                                                                                                                                                                                                                                      RESULT 9
AAY09429
ID AAY0
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Best Local S
Matches 6
                    10-OCT-1997;
                                                     06-OCT-1998;
                                                                                                                          WO9919347-A1
                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                      AAY09429;
                                                                                                                                                                                                                                                                                                                                                        AAY09429 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                Immunoactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 12; Page 155; 183pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rejection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-526397/48.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-APR-1996;
23-SEP-1996;
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                                                                                                                                                                                                                                                                                 14-JUL-1999
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                                                                                                                                                                                            mmune response;
                                                                                                                                                                                                                 mmunoactive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A,
                                                                                                                                                                                                                                                peptide SEQ ID NO:54.
                                                                                                                                                                                                                                                                                   (first entry)
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96SE-00003469.
                    97US-00949024
                                                     98WO-SE001801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eriksson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97WO-SE000574
                                                                                                                                                                                                                                                                                                                                                                                                                                                σ
                                                                                                                                                                                                                                                                                                                                                        peptide; 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 47; DB Pred. No. 2.3;
                                                                                                                                                                                                                                                                                                                                                         Æ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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Nucleic acids encoding cysteine- or methionine-containing peptide(s) which have immuno:stimulatory or immunosuppressive activity - can be to treat, e.g. cancers, infection, auto:immune disease or transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunomodulator; immunosuppressant; immunostimulator; treatment;
transplant rejection; autoimmune disease; cancer; infection.
                                                                                                                                                                                                                                                   Saernstrand
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                                                      used
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The present peptide is an immunosuppressant or immunostimulator. An immunosuppressant can be used to treat transplant rejection or autoimmune disease, e.g. rheumatoid arthritis, systemic lupus erythematosis, Sjogren's syndrome, scleroderma, mixed connective tissue disease, dermatomyositis, polymyositis, Reiter's syndrome, Behcet's disease, type I diabetes, Hashimoto's thyroiditis, Graves' disease, multiple sclerosis, myasthenia gravis, encephalomyelitis, phemphigus vulgaris, vegetans or foliaceus, Senear-Usher syndrome or Brazilian phemphigus. An immunostimulator can be used to treat conditions such as cancer or

2 0; Length 12; Indels <u>,</u>

Gaps

0

immunomodulation; immunosuppression; immunostimulation; se; immunoreactive; autoimmune disease.

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RESULT 10
AAU08670
ID AAU08
XX AAU08
AC AAU08
AC AAU08
AC Thio;
XX Thio;
XX Thio;
XX Cell;
KW cchro;
KW chro;
XX Synt;
XX Synt;
XX Synt;
XX O5-M
PR 07-W
XX O7-W
XX ON-W
PF 05-M
PF 07-W
XX ON-W
XX ON-W
PF 013-S
XX ON-W
PF 05-M
PF 07-W
XX ON-W
PT diff
PT d
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes nucleic acid molecules comprising a coding sequence encoding an immunoreactive peptide and further encoding a protein targeting sequence. The nucleic acid is administered to a patient so that its expression product, an immuno-active peptide, modulates an immune response in a patient. The nucleic acid can also be used to treat cancer, either after surgery to remove a portion of the cancer or after ionizing radiation. A cytokine is also administered in conjunction with the nucleic acid. Cells containing the nucleic acid molecule can also be used for treatment. The immunoactive peptide is immunosuppressive and can be used in patients with autoimmune disease. The present sequence represents a specifically claimed immunoactive peptide from the present
                                                                                          Novel thioredoxin polypeptides and polynucleotides for treating disorder associated with aberrant cellular proliferation or differentiation, e.g., cancer and to identify modulators for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cellular proliferative disorder; differentiative disorder; cancer; carcinoma; sarcoma; tumour; haemopoetic neoplastic disorder; leukaemia; chronic myelogenous leukaemia; cardiovascular disorder; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 32; Page 52; 104pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU08670 standard; protein; 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic genes encoding immunoreactive
                                                                                                                                                                                                                                                                                                                                                                                           07-MAR-2000; 2000US-0187447P
                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-MAR-2001; 2001WO-US007139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-SEP-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thioredoxin domain markov model sequence #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ASTR ) ASTRA
                                                                                                                                                                                                                                                                                                                                   (MILL-) MILLENIUM PHARM INC.
                                                                                                                                                                                                                   2001-589945/66
                                                                                                                                                                                                                                                                              RAJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 WCGPCK 6
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ilarity 100.0%;
Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22105; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eriksson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'n,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lindvall M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 47; DB Pred. No. 2.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             peptides containing cysteine or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Saernstrand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Length 12;
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Disclosure;

Fig 3A; 126pp; English

A 14 kDa molecule designated various Enhancing Factor (ECEF); thioredoxin

names (Eosinophil Cytotoxicity-and ADF) and encoding a polypeptide

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 11
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel human thioredoxin polyperium, fragments of it and allelic variants. 22105 is useful for identifying a compound which modulates the activity or expression of the polypeptide and nucleic acid, where the binds to the polypeptide, and the binding is detected by direct detection of a test compound/polypeptide binding, by using a competition binding assay or an assay for 22105-mediated signal transduction, and the activity of the polypeptide is a redox activity or the ability to modulate protein processing, protein folding and protein secretion. Anti-22105 ana sample. A modulator of 22105 is useful for treating or preventing a disorder such as cellular proliferative or differentiative disorder (e.g. cancers, carcinomas, sarcomas and tumours), haemopoetic neoplastic disorder (e.g. leukaemia and chronic myelogenous leukaemia) on neoplastic disorder (e.g. leukaemia sand chronic myelogenous leukaemia) on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neoplastic disorder (e.g. leukaemia and chronic myelogenous leukaemia) o cardiovascular disorder (e.g. atherosclerosis, ischaemia, thrombosis, hypertension), characterised by aberrant activity or expression of a 22105 nucleic acid or polypeptide in a subject, where the agent is preferably a peptide, a phosphopeptide, a small molecule, an antibody or their combination, or an antisense, a ribozyme, a triple helix molecule, a 22105 nucleic acid or their combination (many examples of each disorde are given in the specification). The present sequence represents a thioredoxin domain sequence derived from a hidden markov model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
04-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 52 AA;
                                                                             Shortened Eosinophil Cytotoxicity-Enhancing Factor proteins - having Eosinophil-stimulating activity, for use in diagnosis and therapy.
                                                                                                                                                                                    Silberstein DS,
                                                                                                                                                                                                                                                        06-APR-1992;
06-JUL-1992;
                                                                                                                                                                                                                                                                                                                                           14-OCT-1993.
                                                                                                                                                                                                                                                                                                                                                                              WO9320107-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                               therapy; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bosinophil cytotoxicity enhancing factor; ECEF; stimulation; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   rECEF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR42822;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR42822 standard; protein;
                                             Example 3; Page 67; 96pp;
                                                                                                                                                     WPI; 1993-336837/42.
                                                                                                                                                                                                                      (BGHM ) BRIGHAM
                                                                                                                                                                                                                                                                                                         06-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 WCGPCK 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 WCGPCK 6
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                                                                                                                                   AAQ49405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first entry)
                                                                                                                                                                                                                                                        92US-00862832
92US-00906842
                                                                                                                                                                                                                                                                                                           93WO-US003310
                                                                                                                                                                                                                      & WOMENS HOSPITAL
                                                                                                                                                                                      Balcewicz-Sablinska MK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 47;
100.0%; Pred. No.
                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 4;
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                                                                                                                                                                                                                                                                             A 14 kDa molecule designated various names (Bosinophil Cytotoxicity-Enhancing Factor (BCEF); thioredoxin and ADF) and encoding a polypeptide of 104 amino acids has been found to support the growth of an EBV transformed B lymphocyte cell line in the manner of IL-1; to induce the expression of IL-2 receptors in a large granular lymphocyte cell line from an ATL patient in the manner of IL-1 and to mediate the growth of the manner of IL-1 and to mediate the growth of the manner of IL-1 and to mediate the growth of the manner of IL-1 and to mediate the growth of the manner of IL-1 and to mediate the growth of the manner of IL-1 and to mediate the growth of the manner of IL-1 and to mediate the growth of the manner of IL-1 and to mediate the growth of the manner of IL-1 and to mediate the growth of the manner of IL-1 and the mediate the growth of the manner of IL-1 and the mediate the growth of the manner of IL-1 and the mediate the growth of the growth of the mediate the growth of the mediate the growth of the growt
because they retain eosinophil stimulating activity but lack proinflammatory dithiol reductase activity. In particular the used for killing tumour cells. (Updated on 25-MAR-2003 to co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shortened Eosinophil Cytotoxicity-Enhancing Factor proteins - having Eosinophil-stimulating activity, for use in diagnosis and therapy.
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04-MAY-1994
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                                                                                                                                           inhibitory properties of interferon gamma. Recombinant ECEF (FECEF) 84 is a truncated version of the 104 amino acid polypeptide comprising the first 84 amino acids of the molecule. Some shortened ECEF's are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; Page 66; 96pp; English.
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06-JUL-1992;
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92US-00906842.
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    they can be correct PN
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RESULT 13
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        can be used in the treatment of various diseases associated with altered mitochondrial function including diabetes mellitus, Huntington's disease, osteoarthritis, Leber's hereditary optic neuropathy (LHON), mitochondrial encephalopathy lactic acidosis and stroke (MELAS), myoclonic epilepsy ragged red fibre syndrome (MERRF) or cancer. Accordingly, these compositions have neuroprotective, nootropic, antidiabetic, anticonvulsant, antiarthritic, osteopathic, ophthalmological and cytostatic activities. This polypeptide sequence is a human heart mitochondrial protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mitochondrial; human; screening assay; diabetes mellitus;
Huntington's disease; osteoarthritis;
Leber's hereditary optic neuropathy; LHON;
mitochondrial encephalopathy lactic acidosis and stroke; MELAS;
myoclonic epilepsy ragged red fibre syndrome; MERRF; cancer;
neuroprotective; nootropic; antidiabetic; anticonvulsant; antiarthritic;
                                                                                                                                                  for therapeutic intervention in treating a disease associated with altered mitochondrial function. Specifically, it refers to a method for identifying proteins of the human heart mitochondrial proteome that are useful for drug screening assays, as well as therapeutic targets. The present invention describes a method for identifying such proteins that
                                                                                                                                                                                                                                                                                                            Identifying a mitochondrial target for drug screening assays and for treating diseases associated with altered mitochondrial function, comprises detecting a modified polypeptide in a sample and correlating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-APR-2002; 2002US-0372843P
17-JUN-2002; 2002US-0389987P
20-SEP-2002; 2002US-0412418P
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                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 312; 180pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human heat mitochondrial protein as a therapeutic target SeqID312
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                                                                                                                                                                                                                                  invention relates to novel mitochondrial targets that can be
                                                                                                                                                                                                                                                                                               the disease
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RESULT 14
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         The present invention describes a method (M1) for producing an oil body CC associated with a recombinant multimeric protein complex (MPC). M1 CC comprises producing in a cell comprising oil bodies a first and second CC comprises producing in a cell comprising oil bodies a first and second CC producing in a cell comprising oil bodies a first and second CC (OB) through an OB-targeting-protein capable of associating with PC (OB) through an OB-targeting-protein capable of associating with OB and CC pl. M1 is useful for producing an oil body associated with a recombinant CC MPC. The oil bodies are further formulated for use in the preparation of CC a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition composition composition composition composition beginned to treat chronic obstructive pulmonary disease (ODD), cataracts, CC diabetes, envenomation, bronchiopulmonary disease (DDD), ulcers, GERD intestinal (GI) bleeding, intestinal bowel disease, passes and nuclear to product product product patential bowel disease, passes (GERD).
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                                                                                                                                                                                                                                                                                                                                                      producing oil body associated with recombinant multimeric protein comple.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
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05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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                                                                                                                                                                                                                                                                                                    Claim 81; Page 243; 362pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-508806/54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SEMB-) SEMBIOSYS GENETICS INC. (SYGN ) SYNGENTA PARTICIPATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rooijen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            c protein; redox protein; thioredoxin; thioredoxin reductase; ophthalmological; antidiabetic; cytostatic; antipsoriatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                G, Deckers l
Zaplachinski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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Moloney M;
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Pred. No.
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G
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RESULT 15
AAB49948
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Matches
                                                Best
                                       Matches
                                                         Query Match
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                                                                                                                                                                                                                                                                                                 17-MAY-1999;
18-MAY-1999;
02-JUN-1999;
                                                                                                                                                                                                                          WPI; 200
N-PSDB;
                                                                                                The present invention provides variants of the Aspergillus oryzae pudisulfide isomerase enzyme. These are capable of reducing disulphide bonds in proteins, which may be the cause of allergies in humans. The proteins can be used to reduce the allergenicity of foods, to treat scleroproteins, in detergents, in food additives and in cosmetics
                                                                                                                                                                                   Protein disulfide isomerase variant having increased reducing properties and decreased redox potential than native proteins, used to reduce allergenicity of allergic proteins in feed, food or cosmetic products.
                                                                                                                                                                                                                                                                                                                                                                                                                         Disulphide bond; allergy; protein disulfide isomerase variant; detergent; food additive; cosmetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A. oryzae protein disulfide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB49948;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 91
                                                                               Sequence
                                                                                                                                                                Claim
                                                                                                                                                                                                                                                         Hjort
                                                                                                                                                                                                                                                                                                                                                                                 WO200070064-A1.
                                                                                                                                                                                                                                                                                                                                                                                                     Aspergillus oryzae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB49948 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABP60964 represent sequence given in the exemplification of the
                                                                                                                                                                                                                                                                                                                                         17-MAY-2000; 2000WO-DK000265
                                                                                                                                                                                                                                                                              (NOVO ) NOVO
                                                  Local
                                                                                                                                                                                                                            2001-070776/08
DB; AAC89030.
                                                                                                                                                                                                                                                                                                                                                                                                                        additive;
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                                                                                                                                                              2; Page 76-77; 82pp; English.
37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 WCGPCK 6
                  1 WCGPCK 6
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                                      Similarity 6; Conserv
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 WCGPCK 42
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                                      100.0%;
ilarity 100.0%;
Conservative 0,
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99DK-00000689.
99US-0137068P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      isomerase
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                                      Score 47; DB Pred. No. 13; Dismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        #3
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12;
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                                                            Length 95
                                        Indels
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                                                                                                                                 s oryzae protein
disulphide
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                                        Gaps
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ABP60779 ID ABP XX

ABP60779 standard; protein; 102

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RESULT 16

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Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a method (M1) for producing an oil body associated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (Pl, P2), where Pl is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and Pl. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                      Sequence 102 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 81; Page 238; 362pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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                                                                                                                                                                                                                                                                                                                                          malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 ABP60964 represent sequence given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-DEC-2001; 2001WO-US050240.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Multimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mycoplasma pneumoniae thioredoxin SEQ
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                                                                                                                                                                                                                                                                                                                      invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SYGN ) SYNGENTA PARTICIPATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rooijen G, Deckers H
Val G, Zaplachinski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           imeric protein; redox protein; thioredoxin; thioredoxin reductase;
body; ophthalmological; antidiabetic; cytostatic; antipsoriatic;
tropic; vulnerary; antibacterial; immunosuppressive; antilcer;
product; milk; wheat; oxidative stress; cataract; diabetes;
29
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6
                                                                                                                                                                 Similarity
                                                            WCGPCK 6
WCGPCK 34
                                                                                                                            Conservative
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                                                                                                                                                           100.0%;
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Moloney M;
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                                                                                                                                                           Score 47; DB
Pred. No. 13;
                                                                                                                            Mismatches
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                                                                                                                                                                                      Length 102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dalmia
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                                                                                                                            Gaps
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RESULT

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Query Match Best Local S Matches 6

Similarity 6; Conser

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Mismatches

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100.0%;

Score 47; Pred. No.

DB 5;

Length 102;

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ABP60755
                              associated with a recombinant multimeric protein complex (MPC). M1
C comprises producing in a cell comprising oil bodies a first and second
c recombinant polypeptide (P1, P2), where P1 is capable of associating with
CC (OB) through an OB-targeting-protein capable of associating with OB to form the MPC and associating the complex with an occlusion body
CC (OB) through an OB-targeting-protein capable of associating with OB and
CP1. M1 is useful for producing an oil body associated with a recombinant
CC MPC. The oil bodies are further formulated for use in the preparation of
CC a food product such as milk or wheat based food product, personal care
CC product which reduces the oxidative stress on the surface area of the
CC human body or used to lighten the skin, or a pharmaceutical composition
CC used to treat chronic obstructive pulmonary disease (COPD), cataracts,
CC diabetes, envenomation, bronchiopulmonary disease, psoriasis,
CC malignancies, reperfusion injury, wound healing, sepsis, gastro
CC (intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD
CC (gastro ossophageal reflux disease). AEN89569 to ABN89593 and ABP60677 to
CC ABP60964 represent sequence given in the exemplification of the present
CC invention. (Updated on 07-AUG-2003 to correct OS field.)
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05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
Sequence
                                                                                                                                                                                                                                                                                                                                            The present invention describes a method (M1) for producing
                                                                                                                                                                                                                                                                                                                                                                                  Claim 81;
                                                                                                                                                                                                                                                                                                                                                                                                                        Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil bodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-DEC-2001; 2001WO-US050240
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06-SEP-2002
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Val G, Zaplachinski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SYNGENTA P
 102
                                                                                                                                                                                                                                                                                                                                                                                  Page 230; 362pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   thioredoxin SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Deckers H, F
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Moloney M;
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**WCGPCK** 

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The present invention describes a method (MI) for producing an oil body cassociated with a recombinant multimeric protein complex (MPC). MI CC comprises producing in a cell comprising oil bodies a first and second crecombinant polypeptide (PI, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (COB) through an OB-targeting-protein capable of associating with OB and CP1. MI is useful for producing an oil body associated with a recombinant CC MPC. The oil bodies are further formulated for use in the preparation of CC a food product such as milk or wheat based food product, personal care CC product which reduces the oxidative stress on the surface area of the CC human body or used to lighten the skin, or a pharmaceutical composition CC malignancies, reperfusion injury, wound healing, sepsis, gastro CC malignancies, reperfusion injury, wound healing, sepsis, gastro CC (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to ABP60964 represent sequence given in the exemplification of the present
Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SEMB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Multimeric oil body; c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chlamydia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Van Rooijen G, Deckers H,
Del Val G, Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gastro
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                                                                       Sequence 102 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Producing oil body associated with recombinant multimeric protein complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-508806/54.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 81; Page 230; 362pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-DEC-2001; 2001WO-US050240.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oesophageal reflux disease.
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                100.0%;
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Moloney M;
Score 47; DB Pred. No. 13; O; Mismatches
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                                 Length 102;
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                                                                                associated with a recombinant multimeric protein complex (MPC). Mi comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (PI, P2), where PI is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and PI. MI is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (CDPD), cataracts, calibrates, reperfusion injury, wound healing, sepsis, gastro intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepbronchiopulmonary disease; malignancy; reperfusion injury; wound he gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                               producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
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05-JUL-2001; 2001US-0302885P
04-DEC-2001; 2001US-00006038
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Del Val G,
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                               (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 ABP60964 represent sequence given in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                           The present invention describes a method (M1) for producing an oil body
                                                                                                                                                                                                                                                                                                                                                                             Claim 81; Page 229; 362pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oil body;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G, Deckers H,
Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     reflux disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            redox protein;
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Moloney M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antidiabetic; cytostatic; antipsoriatic;
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RESULT 20
ABP60777
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         The present invention describes a method (M1) for producing an oil body cassociated with a recombinant multimeric protein complex (MPC). M1 ccomprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition human body or used to lighten the skin, or a pharmaceutical composition capable to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis, capatro cosophageal reflux disease). ABN89595 to ABN89593 and ABP60677 to ABN89594 represent sequence given in the exemplification of the present
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Best Local 9
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n≏l Val G, Zaplachinski
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05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-SEP-2002
                                                                                                                                                                                                                                                                                                                                       Claim 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mycoplasma genitalium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Multimeric protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oesophageal reflux disease.
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                                                                                                                                                                                                                                                                                                                                   Page
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lachinski S,
                                                                                                                                                                                                                                                                                                                                   362pp; English.
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Moloney M;
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Pred. No. 13;
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RESULT 21
ADB09186
ID ADB09186
AX ADB09186
AX ADB09
AX AD
The present invention describes an isolated polynucleotide (I) of CC Alloiococcus otitidis genomic DNA, which encodes an antigenic protein. CC Alloiococcus otitidis is a Gram-positive bacterium. Also described: (1) CC an isolated polypeptide that is encoded by the polynucleotide (I); (2) an isolated polypeptide that is encoded by the polynucleotide (I); its CC complement, degenerate variant or fragment; (3) a genetically engineered CC host cell, transfected, transformed or infected with the vector of (2); (2) an antibody specific for the polypeptide of (1); (5) an immunogenic composition comprising the polypeptide of (1); (5) an immunogenic composition comprising the polypeptide of (1); (5) an immunogenic composition vector; (6) a pharmaceutical composition comprising the expression vector; (6) a pharmaceutical composition comprising an array of the polypeptides of (1), their biological equivalent or fragment; (8) cimmunising against Alloiococcus otitidis by administering to a host the immunogenic composition; (9) detecting and/or identifying Alloiococcus otitidis in the biological sample; (10) a kit comprising a container containing the novel polynucleotide, its degenerate variant or fragment; (6) or the antibody of (4); and (11) producing a polypeptide by culturing the polypeptides of cell under conditions suitable to produce the polynucleotides, polypeptides, antibodies and compositions of the present convention can be used for treating and dagnosing diseases, drug conventions and monitoring of effects during drug clinical trials. The polynucleotides are useful for expressing and detecting Alloiococcus
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Best Local S
Matches 6
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18-NOV-2002; 2002US-0426742P.
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RESULT 22
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ID ABP60
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05-JUL-2001;
04-DEC-2001;
                                                                                  associated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the product which reduces the oxidative stress on the surface area of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                otitidis. The present sequence represents an Alloiococcus otitidis antigen protein from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic, vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-SEP-2002
                               human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts,
                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a method (M1) for producing an oil body
                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 81; Page 227; 362pp;
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Del Val G,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein; redox protein; thioredoxin; thioredoxin
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Zaplachinski S,
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2001US-00006038.
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Moloney M;
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    This invention describes a novel nucleic acid representing a synthetic CC Bax gene. The Bax gene of the invention is useful for identifying Bax-CC resistant yeast or fungi, identifying, or obtaining and identifying CC Candida spp. sequences that are differentially expressed in a pathway CC condida spp. sequences that are differentially expressed in a pathway CC inhibitor sequences of Bax-induced cell death or identifying inhibitors or invention have cytostatic, fungicide, immunosuppressive, virucide and CC vasotropic activity and can be used in vaccines or for gene therapy. The cisolated nucleic acids, polypeptides, pharmaceutical compositions, CC antisense molecules and antibodies are useful as medicaments or in CC preparing a medicament for treating, preventing and/or alleviating CC diseases associated with yeast or fungi or proliferative disorders, such as cancer, or for preventing apoptosis in certain diseases. The compounds CC or polypeptides, or the genetically modified organism are useful for proventing and/orable are useful for treating and/orable are useful for the polypeptides, or the genetically modified organism are useful for treating and/orable are useful for treating and/orable are useful for treating and/orable are useful for the polypeptides.
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04-JAN-2001; 2001EP-00870002
09-JAN-2001; 2001EP-00870003
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                                                                                                                                                                                                                                                                                                                                                                              New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in
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preparing a medicament for modifying the
                                                                                                                                                                                                                                                                                                                                                                certain diseases.
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DB; ABQ76512.
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13;
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RESULT 24
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Matches 6
         This invention describes a novel nucleic acid representing a synthetic Bax gene. The Bax gene of the invention is useful for identifying Bax-resistant yeast or fungi, identifying, or obtaining and identifying Candida spp. sequences that are differentially expressed in a pathway eventually leading to programmed cell death or identifying inhibitors or inhibitor sequences of Bax-induced cell death. The products of the invention have cycostatic, fungicide; immunosuppressive, virucide and vasotropic activity and can be used in vaccines or for gene therapy. The isolated nucleic acids, polypeptides, pharmaceutical compositions, antisense molecules and antibodies are useful as medicaments or in preparing a medicament for treating, preventing and/or alleviating diseases associated with yeast or fungi, preventing and/or alleviating corpolypeptides, or the genetically modified organism are useful for a medicament for treating the compounds or polypeptides, or the genetically modified organism are useful for a medicament for treating the compounds or polypeptides.
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04-JAN-2001; 2001EP-00870002
09-JAN-2001; 2001EP-00870003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            other mammals. The vaccine is useful for immunising against yeast or fungal infections. Apoptosis-related diseases include autoimmune disease, ischaemia, diseases related with viral infections or neurodegenerations. This sequence represents a polypeptide associated with the Bax gene described in the disclosure of the invention
                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Saccharomyces cerevisiae
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                                                                                                                                                                                                                                                                                 Claim
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a medicament for modifying the endogenic flora of humans
                                                                                                                                                                                                                                                                              Fig 1; 344pp; English
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Pred. No.
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RESULT 25
RESULT 25
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This invention describes a novel nucleic acid representing a synthetic Bax gene. The Bax gene of the invention is useful for identifying Bax-cresistant yeast or fungi, identifying, or obtaining and identifying Bax-cresistant yeast or fungi, identifying, or obtaining and identifying CC Candida spp. sequences that are differentially expressed in a pathway eventually leading to programmed cell death or identifying inhibitors or cinhibitor sequences of Bax-induced cell death. The products of the CC invention have cytostatic, fungicide; immunosuppressive, virucide and vasotropic activity and can be used in vaccines or for gene therapy. The isolated nucleic acids, polypeptides, pharmaceutical compositions, CC antisense molecules and antibodies are useful as medicaments or in preparing a medicament for treating, preventing and/or alleviating cd iseasee associated with yeast or fungi or proliferative disorders, such as cancer, or for preventing apoptosis in certain diseases. The compounds or polypeptides, or the genetically modified organism are useful for preparing a medicament for modifying the endogenic flora of humans and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in certain diseases.
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04-JAN-2001; 2001EP-00870002.
09-JAN-2001; 2001EP-00870003.
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Pred. No.
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RESULT
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CC The invention relates to constructing (M1) a strain of diploid fungal cells in which both alleles of a gene are modified, comprising modifying one allele by insertion or replacement by a cassette having an compressible selectable marker and modifying other allele by recombination, of a promoter replacement fragment with a heterologous promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are modified. The diploid fungal cells in which both alleles modified are useful for identifying a gene that is essential to the survival or growth of a fungus, a gene that contributes to the virulence and/or pathogenicity of a fungus, a gene that contributes to the resistance of a diploid fungus to an antifungal agent that inhibits the growth of a diploid fungus and for identifying a therapeutic agent for treatment of a mammalian conspound which modulates the activity of a gene product, preferably enzymatic activity, carbon compound catabolism, biosynthetic, transporter, transcriptional,
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                                                                                                                                                                                                                                                                                                                                   Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inactivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-DEC-2000; 2000US-0259128P.
20-FEB-2001; 2001US-00792024.
22-AUG-2001; 2001US-0314050P.
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signal transduction; DNA replication; cell division; growth;
proliferation; Candida albicans; fungicide; antifungal.
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RESULT 27
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                                                                                                                Dufour GE, Hillman. ...
Dufour GE, Hillman TC, I
Daughtery SC, Dam TC, I
David MH,
Peralta CH, David MH,
Peralta CH, Marwaha R, I
                                                                                                                                                                                                                         28-MAR-2001; 2001US-0279619P.
29-MAR-2001; 2001US-0280067P.
29-MAR-2001; 2001US-0280068P.
16-MAY-2001; 2001US-0291280P.
17-MAY-2001; 2001US-0291280P.
17-MAY-2001; 2001US-0291829P.
19-JUN-2001; 2001US-0291428P.
20-JUN-2001; 2001US-0299776P.
20-JUN-2001; 2001US-0300001P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        translational, signal transduction, DNA replication and cell division activity. The method is useful for identifying a compound having the ability to inhibit growth or proliferation of C. ablicans cells and for treating infection by C. albicans. The present sequence is that of an essential Candida albicans protein used in the method of the invention. Note: The sequence data for this patent is not represented in the printed specification but is based on sequence information supplied to Derwent by the European Patent Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer; cell proliferative disorder; autoimmune disorder; inflammatory disorder; infection; hormonal disorder; metabolic disorder; neurological disorder; astrointestinal disorder; transport disorder; connective tissue disorder; drug screening; proteome analysis; gene therapy; antisense therapy; genetyping; transgenic animal; knock in; disease model; toxicological testing; transcript imaging; electron transfer-associated protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; dithp;
cancer; cell p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 103 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human DITHP electron transfer-associated protein.
                                                                             WPI; 2003-129518/12.
N-PSDB; ACC46545.
                                                                                                                                                                                                                                                                                                                                                               27-MAR-2002; 2002WO-US010056
                                                                                                                                                                                                                                                                                                                                                                                                                  WO200297031-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR41607 standard;
                                                                                                                                                                                                  (INCY-)
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                                                                                                        a CH, Dave u. Marwaha !
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                                                                                                                                                                                                   INCYTE GENOMICS INC.
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                                                                                                                                                                         Jones
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                                                                                                                                             Hillman JL, Yu JY,
C, Dam TC, Liu TF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diagnostic and therapeutic polynucleotide; diagnosis;
                                                                                                                                                                         ĄĮ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein; 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                        Tran AB,
                                                                                                                                  Lewis
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Pred. No.
                                                                                                                      SA,
Lan
                                                                                                                                                                         Dahl CR,
                                                                                                                               Tuason
Nguyen
A, Chen
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                                                                                                                      RY,
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                                                                                                                       Urashka ME;
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                                                                                                                                 Yap PE, Amshey
Kleefeld Y, Ge
Panzer SR, Hai
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                                                                                                                                  Gerstin
Harris B;
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Novel human diagnostic and therapeutic polypeptide useful for ice test compound which specifically binds to a polypeptide encoded diagnostic and therapeutic polynucleotide, and to induce antiboo

identifying ed by human

Claim 27; SEQ ID NO 1142; 591pp; English

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RESULT 28
ADX90669
ID ADX90
XX ADX90
AC ADX90
XX Baker
XX Baker
XX baker
XX Oxido
KW Salin
XX Sacch
XX WO20C
XX O1-JL
XX O7-AL
XX D7-AL
XX BADI;
DR WPI;
DR WPI;
DR WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to novel human diagnostic and therapeutic polynucleotides designated dithp (ACC46094) and to their encoded proteins (DITHP; ABR411136-ABR41812). The invention also relates to proteins (DITHP; ABR41116-ABR41812). The invention also relates to proteins (DITHP; ABR411136-ABR41812). The invention also relates to the dithp cDNA comparison of the invention; recombinant vectors, host cells and transgenic organisms comprising a dithp nucleic acid sequence; the creombinant production of DITHP proteins; antibodies specific for DITHP proteins; microarrays comprising dithp nucleic acid sequences; methods of detecting dithp nucleotide and protein sequences; methods of codetecting dithp nucleotide and protein sequences; methods of codetecting dithp nucleic acid sequences and DITHP proteins may be used in the composition of a wide variety of conditions including cancer and other cell proble Dithp nucleic acid sequences and DITHP proteins may be used in the codingnosis of a wide variety of conditions including cancer and other cell conditions including cancer and other cell conditions including cancer and other cell conditions; fungal or parasitic infections; hormonal disorders; metabolic disorders; neurological disorders gastrointestinal disorders; transport
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Best Local
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   WPI; 2004-226856/21
N-PSDB; ADK90668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Saccharomyces cerevisiae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 baker's yeast; THX2; transgenic; abiotic stress response; crop plant; oxidoreductase stress-related protein; ORSRP; environmental stress; salinity; drought; temperature; chemical; pathogen; glutaredoxin; GRX; thioredoxin; THX; Quantitative Trait Locus; QTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Baker's yeast thioredoxin 2 (THX2) protein SeqID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 103 AA;
                                                                                        Chardonnens A,
                                                                                                                                                                                                              07-AUG-2002; 2002EP-00017671
                                                                                                                                                                                                                                                                       01-JUL-2003; 2003WO-EP006994
                                                                                                                                                                                                                                                                                                                                                                                               WO2004018687-A2
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6; Conserv
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                                                                                                                                                    PLANT SCI GMBH
                                                                                           Puzio P;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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13;
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RESULT 29
ADM32933
ID ADM3293
ADM32933
ID ADM3293
AC ADM32
AC ADM32
AC ADM32
AC AMinc
AC AMIN
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Best Local S
Matches 6
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                                                                                                                                                Increasing the liquefaction of excessively viscous or cohesive mucus sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site reduced state.
                                                                                        Disclosure; Page 64; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Saccharomyces cerevisiae. Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mucus; sputum; thioredoxin; active site; cohesive mucus;
lung disease; cystic fibrosis; enzyme.
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                                                                                                                                                                                                                                                                                                                WPI; 2004-270016/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-SEP-2002;
11-APR-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Amino acid sequence of a thioredoxin polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                         (NAJE-) NAT JEWISH MEDICAL & RES CENT.
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nes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002US-0409960P.
2003US-0462082P.
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The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive

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RESULT 30
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Best Local Similarity
Matches 6; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mucus or sputum. the patient with
A 14 kDa molecule designated various names (Eosinophil Cytotoxicity-Enhancing Factor (ECEF); thioredoxin and ADF) has been found to support the growth of an EBV transformed B lymphocyte cell line in the manner of
                                                                                                                                                                     Shortened Eosinophil Cytotoxicity-Enhancing Factor proteins - havi Eosinophil-stimulating activity, for use in diagnosis and therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-APR-1992;
06-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eosinophil cytotoxicity enhancing factor; ECEF; stimulation; diagnosis; therapy; thioredoxin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR42820
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                                                                                                                   Example 3; Page 65; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-OCT-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                            (BGHM ) BRIGHAM
                                                                                                                                                                                                                                                                         1993-336837/42.
DB; AAQ49403.
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                                                                                                                                                                                                                                                                                                                                                                   DS,
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(first entry)
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92US-00906842.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93WO-US003310
                                                                                                                                                                                                                                                                                                                                                                                                                            & WOMENS HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The method comprises contacting the mucus or sputum of
                                                                                                                                                                                                                                                                                                                                                                Balcewicz-Sablinska MK;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 103;
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RESULT 31
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Matches 6
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Best Local :
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                                                                                 The ADF polypeptide suppresses the release of the pancreatic enzymes amylase and lipase into the blood thereby suppressing pancreatitis. The ADF is also capable of eliminating free radicals which can cause organ damage, as well as refolding denatured proteins which have been denatured by free radicals. The ADF may be isolated from cultured human derived cells and prepared by chemical synthesis or recombinant techniques (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                             Use of a polypeptide having human ADF (thioredoxin) activity - for the prophylaxis or therapeutic treatment of pancreatitis or diseases which accompany pancreatic disorders.
                                                        Sequence 104
                                                                                                                                                                                                    Claim 2; Page 9; 15pp; English.
                                                                                                                                                                                                                                                                                                      WPI; 1994-027626/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADF; thioredoxin; pancreatitis; amylase; lipase; free radicals; protein folding; denaturing; renaturing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human ADF-polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
18-JUL-1994
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                                                                                                                                                                                                                                                                                        N-PSDB; AAQ55111.
                                                                                                                                                                                                                                                                                                                                  Asano T,
                                                                                                                                                                                                                                                                                                                                                                                         19-JUN-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                             (AJIN ) AJINOMOTO KK
                                                                                                                                                                                                                                                                                                                                                                                                                    18-JUN-1993;
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 Similarity 6; Conser
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  Conservative
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(first entry)
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                100.0%;
                                                                                                                                                                                                                                                                                                                                  Isono K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
  0;
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Pred. No.
                Score 47;
Pred. No.
    Mismatches
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                                                                                                                                                                                                                                                                                                                                  Hirakawa T,
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                            Length 104;
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    Indels
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RESULT 32
AAR72389
ID AAR72
XX AAR72
XX AAR72
XX ADF;
XX ACTIV
XX AP17070
XX JP070
PH 20-SE
XX ACTIV
XX AF1;
PH 20-SE
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RESULT 33
AAG80708
ID AAG80
XX AAG80
XX IS-MAG80
XX IS
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Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prepn. of recombinant human ADF by direct expression in E. coli - for treating inflammation and radiation sickness caused by commercially produced free radical(\mathfrak s).
                                                                                                                                                                                                                                                                                                                                S. cerevisiae TRX2 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The DNA sequence encoding human ADF was inserted into a plasmid which was then used to transform E.coli. The E.col was then cultured and the recombinantly produced ADF harvested. Human ADF can be used as a treating agent for inflammation and radiation sickness
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Recombinant human ADF
                                                                                     JP2001288103-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2; Page 7-8; 11pp; Japanese.
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ses 6; Conserv
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DB; AAQ87817.
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                                                                                                                                                                                                                                                   yeast;
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WCGPCK 35
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sickness; fermentation; recombinant.
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                                                                                                                                                                                                                                                   intracellular redox.
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RESULT 34
ABP60927
ID ABP60
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Best Local Similarity
Matches 6; Conser
                                                                                                                                                                                                                                                                                                                                              Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antidicer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis;
                                                                                                                                                                                                                                                                                                        bronchiopulmonary disease; malignancy; gastro intestinal bleeding; intestinal gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes a novel thioredoxin (TRX) composition, stable in solution which contains a yeast TRX derived from Saccharomyces cerevisiae X2180-1A. The TRX composition of the invention can be used for treating and/or preventing lesions and diseases related to intracellular redox conditions. This sequence represents the S. cerevisiae TRX2 protein described in the method of the invention
                                                                                                                                     19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A thioredoxin composition stable in solution used for treating preventing lesions and diseases related to intracellular redox
                        WPI; 2002-508806/54.
                                                                                                                                                                                                                                                   WO200250289-A1
                                                                                                                                                                                                                                                                              Staphylococcus aureus.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Staphylococcus aureus thioredoxin reductase SEQ ID NO:276.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 14-15; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-FEB-2000; 2000JP-00025258
                                                                                              (SEMB-)
                                                                                                                                                                                            19-DEC-2001; 2001WO-US050240
                                                                                                                                                                                                                         27-JUN-2002.
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                                                   Rooijen G, Deckers H,
Val G, Zaplachinski S,
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DB; ABA97065.
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                                                                                              SEMBIOSYS GENETICS INC.
SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard; protein; 104
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                                                    Heifetz PB,
Moloney M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 47; DB Pred. No. 14; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                    reperfusion injury; wound healing;
                                                                  Briggs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                   SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                     Dalmia
                                                                   BK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and/or condition.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                    sepsis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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Producing oil body associated with recombinant multimeric protein complex

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RESULT 35
ABP60803
ID ABP60
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antiuleer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; chronic obstructive pulmonary disease; chronic obstructive pulmona
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil bodies.
                                                                                                                   (SEMB-)
                                                                                                                                                                                                       19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                               Van Rooijen G,
Del Val G, Za
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                                                                                                                                                                                                                                                                                                                              19-DEC-2001; 2001WO-US050240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200250289-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens thioredoxin SEQ ID NO:152
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                                                                                                                   SEMBIOSYS GENETICS INC. SYNGENTA PARTICIPATIONS
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                            G, Deckers H,
Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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100.0%; Pred. No.
                            Heifetz PB,
Moloney M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,</u>
                                                                                                                      AG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB
14;
                                                        Briggs SP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                           Dalmia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 36
ABP60797
ID ABP607
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a method (M1) for producing an oil body associated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with C1 P2 to form the MPC and associating the complex with an occlusion body C2 (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant C2 MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis, malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (GI) bleeding, intestinal bowel disease (IBD), ulcers, GERD (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to ABP60964 represent sequence given in the exemplification of the present
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Best Local S
Matches 6
                                                                                                                                   19-DEC-2000;
05-JUL-2001;
04-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antidicer; food product; milk; wheat; oxidative stress; cataract; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 104
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                                                                                                                                                                                                                                                                                                                                                  27-JUN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bos taurus thioredoxin SEQ ID NO:146.
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                                                                                                                                                                                                                                                                           19-DEC-2001; 2001WO-US050240
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                                   SEMBIOSYS GENETICS INC. SYNGENTA PARTICIPATIONS
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                                                                                                                                       2000US-00742900
2001US-0302885P
2001US-00006038
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 47; DB
Pred. No. 14;
0; Mismatches
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RESULT 37
ABB60812
ID ABB60
XX ABB60812
AC ABB6(
XX ABB6(
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                        oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis, malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (G1) bleeding, intestinal bowel disease (IBD), ulcers, GERD (gastro ossophageal reflux disease). ANN89593 and ABP60677 to
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                                  19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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Del Val G, Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Multimeric protein; redox protein; thioredoxin; thioredoxin reductase;
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6; Conserv
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                                                                                                                                                                    2001WO-US050240
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Moloney M;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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The present invention describes a method (M1) for producing an oil body CC associated with a recombinant multimeric protein complex (MPC). M1 CC comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body CC (D8) through an OB-targeting-protein capable of associating with OB and CP1. M1 is useful for producing an oil body associated with a recombinant CC mfC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition cused to treat chronic obstructive pulmonary disease (COPD), cataracts, and disease, envenomation, bronchiopulmonary disease, psoriasis, malignancies, reperfusion injury, wound healing, sepsis, gastro capable of association of intestinal (G1) bleeding, intestinal bowel disease (IBD), ulcers, GERD capable of association of the surface area of the composition of the surface area of the section of the skin or a pharmaceutical composition considered the section of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
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Sequence 104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Van Rooijen G, Deckers H,
Del Val G, Zaplachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 81; Page 249; 362pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-508806/54.
                                                                                                          ABP60964 represent sequence given in the exemplification of the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Heifetz PB,
Moloney M;
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                                    Query Match
Best Local Similarity
                             Matches
30 WCGPCK 35
             1 WCGPCK 6
                             6;
                              Conservative
                                    100.0%;
                              0,
                                     Score 47;
Pred. No.
                              Mismatches
                                             BB
                                             <u>ب</u>
                             0;
                                            Length 104;
                             0,
                              Gaps
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06-SEP-2002
                                                                                                         ABP60805 standard; protein; 104
 Macaca mulatta
                                   (first entry)
thioredoxin SEQ ID NO:154.
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RESULT 38
ABP60805
ID ABP60
XX ABP60
XX ABP60
XX ABP60
XX ABP60
XX ABP60
XX MACA
XX MALT
KW MILT
KW OILL
KW VASC
KW Chrol
KW SAB
KW GAB
XX GAB
XX MAC
XX PD 27
XX PD 27
XX 19 chronic obstructive pulmonary disease; envenomation; psoriasis; bronchiopulmonary disease; malignancy; reperfusion injury; wound gastro intestinal bleeding; intestinal bowel disease; ulcer; Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antidicer; food product; milk; wheat; oxidative stress; cataract; diabetes; 19-DEC-2000; 2000US-00742900 19-DEC-2001; 2001WO-US050240 Macaca mulatta. gastro oesophageal disease healing;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a method (M1) for producing an oil body associated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal caree product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease (COPD), cataracts, analignancies, reperfusion injury, wound healing, sepsis, gastro (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to ABP80964 represent sequence given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil bodies.
                                                                                                                          chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUL-2001; 2001US-0302885P. 04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                          Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antidicer; food product; milk; wheat; oxidative stress; cataract; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 81; Page 247; 362pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Van Rooijen G, Deckers H,
Del Val G, Zaplachinski S,
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                                                        WO200250289-A1
                                                                                         Mus musculus.
                                                                                                                                                                                                                                                                                  Mus musculus thioredoxin SEQ ID NO:155.
                                                                                                                                                                                                                                                                                                                      06-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                        ABP60806 standard; protein; 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity nes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                     (first entry
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Moloney M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 47; DB 5; Length 104; Pred. No. 14;
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                                                                                                                                                                                                                                                                                                                           RESULT 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 6
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e.g. redox proteins and immunoglobulins comprises producing recombinant
polypeptides capable of forming the complex in cells comprising oil
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05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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Del Val G, Zaplachinski
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 104
                                                                                                                                                                                                                                                                                            ABP60809 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         81; Page 247; 362pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                       1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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lachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
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Moloney M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 47; DB
Pred. No. 14;
0; Mismatches
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Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immonsuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing;

Oryctolagus cuniculus thioredoxin SEQ ID NO:158.

(first entry)

bronchiopulmonary disease; malignancy; reperi gastro intestinal bleeding; intestinal bowel gastro oesophageal reflux disease.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
         Multimeric protein; redox protein; thioredoxin; thioredoxin reductable; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antilcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 104 AA;
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05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
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                                                                                                                                                                           Rattus norvegicus thioredoxin SEQ ID NO:159.
                                                                                                                                                                                                              06-SEP-2002
                                                                                                                                                                                                                                                  ABP60810
                                                                                                                                                                                                                                                                                  ABP60810 standard; protein; 104 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rooijen G, Deckers I
Val G, Zaplachinski
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                                                                                                                                                                                                                                                                                                                                                                                                     1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                  6; Conserv
                                                                                                                                                                                                                                                                                                                                                                              WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                              (first entry)
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lachinski S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 47;
100.0%; Pred. No.
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Moloney M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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RESULT 42
ABP60801
ID ABP60
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                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 6
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05-JUL-2001; 2001US-0302885P
04-DEC-2001; 2001US-00006038
vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis: sens; bronchiopulmonary disease;
                                                           Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic;
                                                                                                              Gallus gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 81;
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Del
                                                                                                                                                  06-SEP-2002
                                                                                                                                                                               ABP60801;
                                                                                                                                                                                                                 ABP60801 standard;
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Val G,
                                                                                                                                                                                                                                                                                                    30 WCGPCK 35
                                                                                                                                                                                                                                                                                                                                   1 WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                       104
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                                                                                                                                                                                                                                                                                                                                                                   100.0%; ilarity 100.0%; Conservative
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Zaplachinski S,
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                                                                                                                                                (first
                                                                                                                thioredoxin SEQ ID NO:150.
                                                                                                                                                                                                                 protein; 104 AA
                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Heifetz PB,
Moloney M;
                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                   Score 47; DB Pred. No. 14; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a
                                                                                                                                                                                                                                                                                                                                                                                        14;
                                                                                                                                                                                                                                                                                                                                                                                                      DB
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                                                                                                                                                                                                                                                                                                                                                                                                    Length 104;
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                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dalmia
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0,

bronchiopulmonary disease;

wound healing;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC associated with a recombinant multimeric protein complex (MPC). M1
CC comprises producing in a cell comprising oil bodies a first and second
CC recombinant polypeptide (P1, P2), where P1 is capable of associating with
CC P2 to form the MPC and associating the complex with an occlusion body
CC (OB) through an OB-targeting-protein capable of associating with OB and
CC P1. M1 is useful for producing an oil body associating with a recombinant
CC MPC. The oil bodies are further formulated for use in the preparation of
CC a food product such as milk or wheat based food product, personal care
CC product which reduces the oxidative stress on the surface area of the
CC human body or used to lighten the skin, or a pharmaceutical composition
CC used to treat chronic obstructive pulmonary disease (COPD), cataracts,
CC diabetes, envenomation, bronchiopulmonary disease, psoriasis,
CC malignancies, reperfusion injury, wound healing, sepsis, gastro
CC (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to
CC intestinal (GI) bleeding, intestinal bowel disease (TBD), ulcers, GERD
CC (gastro esophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to
CC intestinal sequence given in the exemplification of the present
                                                                                                                                                                                               ABG93115
                                                                                                                                                                                                                RESULT 43
                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gastro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gastro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200250289-A1
                                                                    S. cerevisiae BAX-associated protein fragment SEQ ID 188.
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 81; Page 245-246; 362pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-508806/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-DEC-2001; 2001WO-US050240
vasotropic; vaccine; gene therapy; proliferative disorder; cancer;
apoptosis; fungal; yeast; infection; autoimmune disease; ischaemia;
                                   Bax; Bax-resistance;
                                                                                                          21-NOV-2002
                                                                                                                                                                             ABG93115 standard; protein; 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SEMB-) SEMBIOSYS GENETICS INC. (SYGN ) SYNGENTA PARTICIPATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rooijen G, Deckur
                                                                                                                                                                                                                                                                        30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 intestinal bleeding; intestinal bowel disease; ulcer; oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gallus.
                                                                                                                                                                                                                                                                                                          1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                           Similarity 6; Conserv
                                                                                                                                                                                                                                                                      WCGPCK 35
                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                          (first entry)
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lachinski S,
                cytostatic; fungicide; immunosuppressive; virucide; gene therapy; proliferative disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 47;
100.0%; Pred. No.
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Moloney M;
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                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                               DB 5; Length 104;
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ADK90667 standard;

protein;

0

baker's yeast; THX1; transgenic; abiotic stress response; crop plant; oxidoreductase stress-related protein; ORSRP; environmental stress; salinity; drought; temperature; chemical; pathogen; glutaredoxin; GRX

Baker's yeast thioredoxin 1 (THX1) protein SeqID 12

(first

entry)

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RESULT 44
ADK90667
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                                                                                                   S
                                                                                                                                                                                                                                    CC Bax gene. The Bax gene of the invention is useful for identifying Bax-
CC resistant yeast or fungi, identifying, or obtaining and identifying
CC candida spp. sequences that are differentially expressed in a pathway
CC eventually leading to programmed cell death or identifying inhibitors or
CC inhibitor sequences of Bax-induced cell death. The products of the
CC invention have cytostatic, fungicide; immunosuppressive, virucide and
CC vasotropic activity and can be used in vaccines or for gene therapy. The
CC intisense molecules and antibodies are useful as medicaments or in
CC preparing a medicament for treating, preventing and/or alleviating
CC diseases associated with yeast or fungi or proliferative disorders, such
CC or polypeptides, or the genetically modified organism are useful for
CC preparing a medicament for modifying the endogenic flora of humans and
CC or polypeptides. The vaccine is useful for immunising against yeast or
CC fungal infections. Apoptosis related diseases include autoimmune disease,
CC ischaemia, diseases related with viral infections or neurodegenerations.
CC This sequence represents a polypeptide associated with the Bax gene
CC described in the disclosure of the invention
                                                                                                                                                    Query Match
Best Local S
                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-DEC-2000; 2000EP-00870318
04-JAN-2001; 2001EP-00870002
09-JAN-2001; 2001EP-00870003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           certain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Saccharomyces cerevisiae.
                                                                                                                                                                                                         Sequence 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 36; Fig 1; 344pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contreras
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neurodegeneration; cell death
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (JANC ) JANSSEN PHARM NV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002-667002/71.
DB; ABQ76381.
                                                                     30
                                                                                                       1 WCGPCK 6
                                                                                                                                       Similarity 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diseases.
                                                                                                                                         Conservative
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A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         describes a novel nucleic acid representing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eberhardt
                                                                                                                                                       100.0%;
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                                                                                                                                                         Score 47;
Pred. No.
                                                                                                                                         Mismatches
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                                                                                                                                                                           <u>ب</u>
                                                                                                                                         0
                                                                                                                                                                           Length 104;
                                                                                                                                         Indels
                                                                                                                                         0,
                                                                                                                                         Gaps
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ADC59801
ADC59901
XX
ADC599
AC ADC59
XX
I5-JU
XX
Chip;
KW chip;
KW cspB;
KW ldh;
KW ldh;
XX
Bacil
NX
Bacil
PN DE102
XX
I1-SE
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to novel transgenic plant cells transformed by genes encoding proteins associated with the abiotic stress response. Specifically, it refers to transgenic crop plants expressing the heterologous oxidoreductase stress-related protein (ORSRP) that confers increased tolerance and/ or resistance to environmental stresses such as salinity, drought, temperature, chemicals or pathogens. The present invention describes the ORSRPs as heat-stable glutaredoxin (GRX) and/ or thioredoxin (THX) proteins, where expression is under the control of an inducible tissue-specific or developmentally-specific promoter and furthermore it provides antisense oligos and RNA interference molecules to inhibit ORSRP expression. In addition, these nucleic acids are useful as Quantitative Trait Locus (QTL) markers that can map genetic loci associated with environmental stress tolerance. This polypeptide sequence
                                                                                                                                       cspB; des; dnaK; eno; glnR; groEL; groL; ibpA; ibpB; katA; katE; lctP; ldh; opuAB; phoA; phoD; pstS; purC; purN; pyrB; pyrP; sigB; tnrA; trxA; ydjF; fermentation monitoring; process control; increased efficiency.
                                                                                                                                                                                                                            B. subtilis metabolic pathway gene, trxA, protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 104 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New transgenic plant cell transformed by oxidoreductase stress-related protein (ORSRP) coding nucleic acid, useful for producing a transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-AUG-2002; 2002EP-00017671
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Saccharomyces cerevisiae
                                                                                                        Bacillus subtilis.
                                                                                                                                                                                             chip; metabolic pathway; acoA; ahpC; ahpF; citB; clpC; clpP; codY; cspA;
                                                                                                                                                                                                                                                                   15-JUL-2004
                                                                                                                                                                                                                                                                                                                                   ADO59801 standard; protein; 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; SEQ ID NO 12; 140pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plant with increased environmental stress tolerance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; ADK90666.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chardonnens A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  thioredoxin; THX; Quantitative Trait Locus; QTL
 11-SEP-2002; 2002DE-01042433.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        is a baker's yeast THX protein of the invention.
                                    25-MAR-2004
                                                                     DE10242433-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BADI ) BASF PLANT SCI GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2004-226856/21.
                                                                                                                                                                                                                                                                                                                                                                                                                             30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                             WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                WCGPCK 6
                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Puzio P;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 47; DB Pred. No. 14; O; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8; Length 104;
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RESULT 46
ADS88157
ID ADS88
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a chip that carries probes for at least one regulated genes involved in the corresponding metabolic pathways in other organisms. The specified genes are acoA, ahpC, ahpF, citBC, clpP, codY, cspA, cspB, des, dnaK, eno, glnR, groEL, groL, ibpA, ibpB, katA, katE, lctP, ldh, opuAB, phoA, phoD, pstS, purC, purN, pyrB, pyrP, sigB, tnrA, trxA and ydjF. The specification includes a table identifying the products of these genes. The chips are used to determine the physiological condition of organisms being used in a biological process, especially for monitoring fermentations for production of a protein or low molecular weight compound, but also for laboratory-scale cloning and expression analysis. The method is quicker than gel electrophoretic methods currently used, so allows a more rapid intervention to process control and thus increased efficiency. The present sequence represents the amino acid sequence of a B. subtilis metabolic pathway gene protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chip carrying probes for specific genes, useful for rapid monitoring organism status, particularly during fermentation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-271050/26.
N-PSDB; ADO59800.
                                           26-SEP-2002;
10-FEB-2003;
                                                                                                                                                          WO2004035783-A2
                                                                                                                                                                                        Homo
                                                                                                                                                                                                                       antirheumatic;
                                                                                                                                                                                                                                    TNP-alpha; chronic inflammatory disease; rheumatoid arthritis; inflammatory bowel disease; infectious disease; septic shock; bacterial infection; neurological disease; stroke-induced inflammation; neurodegenerative disease; cancer; antiinflammatory; antiarthritic;
                                                                                                                                                                                                                                                                                                                                  Human protein of a TNF-alpha signalling pathway protein complex SeqID
                                                                                                                                                                                                                                                                                                                                                                    18-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                  ADS88157 standard; protein; 104 AA
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Voigt B;
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                                                                                          24-SEP-2003; 2003WO-EP050655
                                                                                                                          29-APR-2004
                                                                                                                                                                                                                                                                                   protein complex; tumour necrosis factor-alpha signalling pathway; TNF-alpha; chronic inflammatory disease; rheumatoid arthritis;
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6; Conserva
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UNIV GREIFSWALD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 104 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
ilarity 100.0%;
Conservative 0
                                          2002EP-00021809.
2003EP-00100274.
                                                                                                                                                                                                                        cytostatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 47; DB
Pred. No. 14;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schweder T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8; Length 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hecker M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Juergen
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(CELL-) CELLZOME AG

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RESULT 47
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                present invention describes a protein complex and kit that are useful for diagnosing, prognosing or treating chronic inflammatory diseases such as rheumatoid arthritis and inflammatory bowel disease; infectious diseases such as septic shock and bacterial infections; neurological diseases such as stroke-induced inflammation in neurons; neurological diseases such as stroke-induced inflammation in neurons; neurodegenerative diseases such as troke-induced inflammation in neurons; neurodegenerative diseases and cancer. Accordingly, these complexes can be used for the development of pharmaceutical compositions that exhibit antiinflammatory, antiarthritic, antirheumatic, cytostatic and antibacterial activities and can be used for gene therapy purposes. In particular, the invention further provides siRNA-oligonucleotides useful for inhibiting protein expression for in vitro or cell culture assays. This polypeptide is a human protein that can be used in combination with other proteins provided in the specification to form novel complexes of the TNF-alpha signalling pathway of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to novel protein complexes of the tumour necrosis factor-alpha (TNF-alpha) signally, it refers to methods for preparing these complexes comprising at least two component proteins, as well as screening methods to identify modulators of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New protein complex comprising at least one first and second protein the Tumor Necrosis Factor-alpha(TNF-alpha)-signaling pathway, useful diagnosing or treating inflammation, neurological diseases, infectiou diseases or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bouwmeester T, I
Superti-Furga G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example;
                                  12-JUN-1987;
31-MAY-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 104 AA;
                                                                                                                                                                    EP299206-A.
                                                                                                                                                                                                                                                                                                                                      Human adult T cell leukaemia
                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-2003
25-JUN-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAP92141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAP92141 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pathway,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proteins,
(AJIN ) AJINOMOTO KK
                                                                                          10-JUN-1988;
                                                                                                                               18-JAN-1989.
                                                                                                                                                                                                                        Region
                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                   immunodeficiency disease
                                                                                                                                                                                                                                                                                                                                                                         Recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30 WCGPCK 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              which include antibodies, agonists and antagonists thereof.
                                                                                                                                                                                                                                                                                                                                                                         human adult T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ħ
                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
(first entry)
                                  87JP-00146348.
88JP-00134218.
                                                                                          88EP-00109311
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NO 12; 1980pp;
                                                                                                                                                                                                                      Location/Qualifiers
2. .105
                                                                                                                                                                                                         /note= "Claim 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kruse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₫
                                                                                                                                                                                                                                                                                                                                                                           cell leukaemia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                      derived factor; hADF; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Þ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB
14;
                                                                                                                                                                                                                                                                                                                                                                           derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ŧ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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RESULT 48
AAR42819
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        A 14 kDa molecule was isolated by three separate groups and found to support the growth of an EBV transformed B lymphocyte cell line in the manner of IL-1. It was also found to induce the expression of IL-2 receptors in a large granular lymphocyte cell line from an ATL patient in the manner of IL-1. The molecule has been designated various names, these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA encoding the polypeptide was sequenced from a gene bank prepd. from mRNA isolated from ATL-2 cells from patient with adult T leukaemia virus. Vectors contg. the DNA can be used to transform host cells for prodn. of hADF polypeptide. The polypeptide causes differentiation and induces growth of lymphocytes and fibroblasts. The N-terminal Met is optional. Similar peptides with deletions, sustns. and allelic derivs. may also be used. See also AAP94809. (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                     Shortened Eosinophil Cytotoxicity-Enhancing Eosinophil-stimulating activity, for use in
                                                                                                                                                                                                                         06-APR-1992;
06-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2003
04-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 105 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Recombinant human adult T used for treating cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yodoi J,
                                                                                                                                                                              Silberstein DS,
                                                                                                                                                                                                                                                              06-APR-1993;
                                                                                                                                                                                                                                                                                                                                                        therapy; thioredoxin
                                                                                                                                                                                                                                                                                                                                                                  Eosinophil
                                                                                                                                                                                                                                                                                                                                                                                            ECEF/thioredoxin/ADF
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR42819;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR42819 standard;
                                                                                Disclosure; Fig
                                                                                                                                                                                                                                                                                     14-OCT-1993
                                                                                                                                                                                                                                                                                                            WO9320107-A1
                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                    (BGHM ) BRIGHAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                      1993-336837/42.
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Eosinophil Cytotoxicity-Enhancing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tagaya Y,
                                                                                                                                                                                                                                                                                                                                                                  cytotoxicity enhancing factor; ECEF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
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92US-00906842.
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و
                                                                                                                                                                                                     & WOMENS HOSPITAL
                                                                                                                                                                            Balcewicz-Sablinska
                                                                             96pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .0%;
                                                                                English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell leukaemia derived factor polypeptide immuno-deficiency disease etc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 47;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ₽
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Factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB
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                                                                                                       Factor proteins diagnosis and th
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                                                                                                                                                                                                                                                                                                                                                                      stimulation;
(ECEF)
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                                                                                                       therapy.
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                                                                                                                                                                                                                                                                                                                                                                      diagnosis;
                                                                                                                  having
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RESULT 50
AAR47856
ID AAR47
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AAR37700
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Matches
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Best Local Similarity
                                                                                                                 Matches
                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                    New human
prepn. of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADF. The molecule has also been found to mediate the growth inhibitory properties of interferon gamma. Some shorteened ECEF's are useful because they retain eosinophil stimulating activity but lack proinflammatory dithiol reductase activity. In particular they can be used for killing tumour cells. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                              Sequence 105 AA;
                                                                                                                                                                                 This sequence represents human ADF. The DNA encoding this sequence may used in the production of a transgenic mouse. The transgenic mouse contains, in the 5' to 3' direction, a human beta-actin promoter, the human ADF coding gene, a termination codon, a human beta-actin poly(A) signal and a human beta-actin terminator. The transgenic mouse is resistant to stress. It is useful in the research of the mechanism of stress and for the development of anti-inflammatory drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JP05130819-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR37700 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 105 AA;
            AAR47856 standard;
                                                                                                                                                                                                                                                                             Claim 3; Page 5; 10pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                     12-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                            28-MAY-1993.
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                                                                                                                                                                                                                                                                                                                                                                                              12-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               resistant; stress;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                                                         (AJIN ) AJINOMOTO KK
                                                                                                                Local Similarity
les 6; Conserv
                                                                                                                                                                                                                                                                                                                                                  1993-208254/26.
                                                                   31
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                                                                                          μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADF; transgenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6,
                                                                                          WCGPCK 6
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                                                                    WCGPCK 36
                                                                                                                                                                                                                                                                                                 ADF transgenic mouse - is resistant to stress and is useful antiinflammatory drugs.
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                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                       91JP-00295618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               enic; mouse; beta-actin; promoter; terminator;
anti-inflammatory drugs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein; 105
          protein; 105
                                                                                                                           100.0%;
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Pred. No.
                                                                                                                            Score 47;
Pred. No.
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                                                                                                                 Mismatches
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14;
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                                                                                                                                       Length 105;
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                                                                                                                 Indels
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                                                                                                                 0,
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                                                                                                      Gaps
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RESULT 51
AAR72390
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                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                  The ADF polypeptide suppresses the release of the pancreatic enzymes amylase and lipase into the blood thereby suppressing pancreatitis. The ADF is also capable of eliminating free radicals which can cause organ damage, as well as refolding denatured proteins which have been denatured by free radicals. The ADF may be isolated from cultured human derived cells and prepared by chemical synthesis or recombinant techniques (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Use of a polypeptide having human ADF (thioredoxin) activity - for the prophylaxis or therapeutic treatment of pancreatitis or diseases which accompany pancreatic disorders.
          28-MAR-1995.
                              JP07079780-A.
                                                                                                                          Recombinant human
                                                                                                                                               10-NOV-1995
                                                                                                                                                                     AAR72390;
                                                                                                                                                                                       AAR72390 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 2; Page 12; 15pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Asano T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-JUN-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF; thioredoxin; pancreatitis; amylase; lipase; free radicals; protein folding; denaturing; renaturing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human ADF-polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
18-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR47856;
                                                                                  Homo
                                                                                                      ADF; inflammation; radiation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JAN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                  ctive-site
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                                                                                 sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1994-027626/04.
                                                                                                                                                                                                                                            31
                                                                                                                                                                                                                                                                 μ
                                                                                                                                                                                                                                             WCGPCK 36
                                                                                                                                                                                                                                                                WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kenmochi T,
                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92JP-00161454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93EP-00109794.
                                                  Location/Qualifiers 32. .35
                                                                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isono
                                                                                                                                                                                          105 AA
                                                                                                                                                                                                                                                                                      0;
                                                                                                      sickness;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ~
                                                                                                                                                                                                                                                                                                 Score 47; DB
Pred. No. 14;
                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hirakawa
                                                                                                      fermentation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ,
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                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hamuro
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                                                                                                                                                                                                                                                                                      Indels
                                                                                                      recombinant
                                                                                                                                                                                                                                                                                      0;
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RESULT 52
AAW69567
ID AAW69
XX AAW69
AC AAW69
XX AAW69
AC AAW69
XX Human
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Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human;
Ref-1.
New thioredoxin variants with amino acid substitutions for cysteine except at active centre for increasing stability in non-reducing conditions - used with Ref-1 to enhance AP-1 transcriptional activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The DNA sequence encoding human ADF was inserted into a plasmid which was then used to transform E.coli. The E.coli was then cultured and the recombinantly produced ADF harvested. Human ADF can be used as a treating agent for inflammation and radiation sickness
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW69567 standard; protein; 105
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                                                                                        WPI; 1998-364621/32.
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nes 6; Conserv
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Pred. No.
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14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a human thioredoxin (TRX) variant derived from the wild-type sequence (obtained from Genbank), as stated in the specification. TRX variants can be made stable under non-reducing conditions by substituting at least one or all Cys residues by other amino acid residues, except that Cys residues in the active centre of TRX remain unmodified. Also described in the present invention is a factor for enhancing the transcriptional activity of AP-1 which comprises the TRX variants can be used with Ref-1 to enhance AP-1 transcriptional activity. TRX variants do not form multimers and are stable under non-reducing
                                               This invention describes novel antisense oligonucleotides against thioredoxin and thioredoxin reductase gene which have cytostatic activity and are useful for inhibiting tumour growth and metastasis in mammals. They may also be used as hybridization probes to detect the presence of the thioredoxin and thioredoxin reductase mRNAs in mammalian cells. They may also be used as molecular weight markers. The antisense oligonucleotides are nuclease resistant due to the presence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thioredoxin; thioredoxin reductase; l cytostatic; tumour growth inhibitor; phosphorothioate linkage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 105
                                                                                                                                                                                     genes,
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                                                                                                                                                            Disclosure;
                                                                                                                                                                                                 Antisense oligonucleotides against thioredoxin and thioredoxin reductase
                                                                                                                                                                                                                                                                                                                        30-JAN-1998;
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                                                                                                                                                                                     useful
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                                                                                                                                                                                   inhibiting tumor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     reductase; human; antisense; primer; metastasis;
h inhibitor; detection; nuclease resistant;
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Pred. No.
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Sequence 105

B

phosphorothicate

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linkages.

This sequence

represents

protein

Length 105;

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RESULT 54
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                                               Query Match
Best Local S
Matches 6
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Best Local Similarity
Matches 6; Conser
                                                                                                                   by the street from mRNAs encoding secreted proteins. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion vectors
                                                                                                  Sequence 105 AA;
                                                                                                                                                                                                                                                                                                           Claim 13; SEQ ID NO 8037; 71pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                   New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures.
                                                                                                                                                                                                                                                                                                                                                                                                                                Dumas Milne Edwards
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                                               6; Conserv
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                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chromosome
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                                                                                                                                                                                                                                                                                                                                                                                                                                Duclert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence tag; secreted protein; cDNA isolation; mapping.
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                                               Score 47; DB:
Pred. No. 14;
0; Mismatches
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RESULT 55 ABP60700

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31

WCGPCK 6 WCGPCK 36

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Indels

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The present invention describes a method (M1) for producing an oil body CC associated with a recombinant multimeric protein complex (MPC). M1 CC comprises producing in a cell comprising oil bodies a first and second CC recombinant polypeptide (P1, P2), where P1 is capable of associating with CC P2 to form the MPC and associating the complex with an occlusion body CC (OB) through an OB-targeting-protein capable of associating with OB and CC III. M1 is useful for producing an oil body associated with a recombinant CC MPC. The oil bodies are further formulated for use in the preparation of CC a food product such as milk or wheat based food product, personal care CC product which reduces the oxidative stress on the surface area of the Numan body or used to lighten the skin, or a pharmaceutical composition CC diabetes, envenomation, bronchiopulmonary disease (COPD), cataracts, CC diabetes, envenomation, bronchiopulmonary disease (COPD), cataracts, CC diabetes, envenomation, bronchiopulmonary disease (IBD), ulcers, GERD CC (gastro ossophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to (manipulation of the present in the exemplification of the present
Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-DEC-2000;
05-JUL-2001;
04-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic; vasotropic; vulnerary; antibacterial; immunosuppressive; antiuleer; food product; milk; wheat; oxidative stress; cataract; diabetes; chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer; gastro oesophageal reflux disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil bodies.
                                                                        Sequence 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 203; 362pp; English.
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Val G, Zaplachinski S,
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100.0%; Score 47; DB ilarity 100.0%; Pred. No. 14; Conservative 0; Mismatches
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2001US-0302885P.
2001US-00006038.
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Moloney M;
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                                     ŝ
                                       Length 105
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RESULT 56
ABP60757
                                                                       The present invention describes a method (M1) for producing an oil body cascoiated with a recombinant multimeric protein complex (MPC). M1 comprises producing in a cell comprising oil bodies a first and second recombinant polypeptide (P1, P2), where P1 is capable of associating with P2 to form the MPC and associating the complex with an occlusion body (OB) through an OB-targeting-protein capable of associating with OB and P1. M1 is useful for producing an oil body associated with a recombinant MPC. The oil bodies are further formulated for use in the preparation of a food product such as milk or wheat based food product, personal care product which reduces the oxidative stress on the surface area of the human body or used to lighten the skin, or a pharmaceutical composition used to treat chronic obstructive pulmonary disease (COPD), cataracts, diabetes, envenomation, bronchiopulmonary disease, psoriasis, malignancies, reperfusion injury, wound healing, sepsis, gastro intestinal (G1) bleeding, intestinal bowel disease (IBD), ulcers, GERD (gastro oesophageal reflux disease). ABN89569 to ABN89593 and ABP60677 to ABP60964 represent sequence given in the exemplification of the present
Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Producing oil body associated with recombinant multimeric protein complex e.g. redox proteins and immunoglobulins comprises producing recombinant polypeptides capable of forming the complex in cells comprising oil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-DEC-2000; 2000US-00742900.
05-JUL-2001; 2001US-0302885P.
04-DEC-2001; 2001US-00006038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chronic obstructive pulmonary disease; envenomation; psoriasis; sepsis; bronchiopulmonary disease; malignancy; reperfusion injury; wound healing; gastro intestinal bleeding; intestinal bowel disease; ulcer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Multimeric protein; redox protein; thioredoxin; thioredoxin reductase; oil body; ophthalmological; antidiabetic; cytostatic; antipsoriatic;
                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Van Rooijen
Del Val G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gastro oesophageal reflux disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 81; Page 230; 362pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-508806/54.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vasotropic; vulnerary; antibacterial; immunosuppressive; antiulcy
food product; milk; wheat; oxidative stress; cataract; diabetes;
                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SYNGENTA PARTICIPATIONS
 6,
                                                                         105
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   Conservative
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                  100.0%;
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Moloney M;
 0
                Score 47;
Pred. No.
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   Mismatches
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                                 Length 105
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Query Match Best Local Similarity 6; Conserve

Conservative

100.0%;

.0%; Score 47; .0%; Pred. No. 0

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5.

Length 105;

Mismatches

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Gaps

0,

Sequence 105

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RESULT 57
ABG93363
ID ABG93
AC ABG92
XX ABG93
XX ABG92
XX BAX;
WW BAX;
WW BAX;
WW ABBG
XX BAX;
WW ABBG
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WW ABBG
XX BAX;
WW ABGG
XX BAX;
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                                                                  isolated nucleic acids, polypeptides, pharmaceutical compositions, antisense molecules and antibodies are useful as medicaments or in preparing a medicament for treating, preventing and/or alleviating diseases associated with yeast or fungi or proliferative disorders, such as cancer, or for preventing apoptosis in certain diseases. The compounds or polypeptides, or the genetically modified organism are useful for preparing a medicament for modifying the endogenic flora of humans and other mammals. The vaccine is useful for immunising against yeast or fungal infections. Apoptosis-related diseases include autoimmune disease, ischaemia, diseases related with viral infections or neurodegenerations. This sequence represents a polypeptide associated with the Bax gene described in the disclosure of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel nucleic acid representing a synthetic Bax gene. The Bax gene of the invention is useful for identifying Bax-resistant yeast or fungi, identifying, or obtaining and identifying Candida spp. sequences that are differentially expressed in a pathway eventually leading to programmed cell death or identifying inhibitors inhibitor sequences of Bax-induced cell death. The products of the invention have cytostatic, fungicide; immunosuppressive, viricide and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated nucleic acid representing a synthetic BAX-gene, useful as medicament for treating, preventing and/or alleviating yeast or fungal infections or proliferative disorders, or for preventing apoptosis in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-DEC-2000; 2000BP-00870318
04-JAN-2001; 2001BP-00870002
09-JAN-2001; 2001BP-00870003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vasotropic; vaccine; gene therapy; proliferative disorder; cancer; apoptosis; fungal; yeast; infection; autoimmune disease; ischaemia; neurodegeneration; cell death.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bax; Bax-resistance; cytostatic; fungicide; immunosuppressive; virucide; vasotropic; vaccine; gene therapy; proliferative disorder; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-DEC-2001; 2001WO-EP015398
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vasotropic activity and can be used in vaccines or for gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (JANC ) JANSSEN PHARM NV.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            36; Fig 2; 344pp; English.
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WCGPCK 36 WCGPCK 6

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RESULT 58
ABO07225
The invention relates to identifying (M1) a candidate p53 pathway CC modulating agent, by contacting an assay system comprising a purified HM CC polypeptide (human orthologue of genes that modify the p53 pathway in CC Drosophila) or nucleic acid with a test agent under conditions, where but CC for the presence of the test agent, the system provides a reference cativity, and detecting a test agent, the system provides a reference contacting a cell defective in p53 function with a candidate modulator CC that specifically binds to a HM polypeptide comprising an HM amino acid contacting a cell (comprising contacting the cell with an agent that specifically binds to a HM polypeptide comprising (M3) a p53 pathway in a mammalian cell (comprising contacting the cell with an agent that conspective in a patient (comprising the sample with a probe for HM comparison indicates a likelihood disease). (M1) is useful for identifying modulators of the p53 pathway. A probe for HM comparison indicates a likelihood disease). (M1) is useful for diagnosing breast, colon, kidney, lung and ovarian cancer, in a patient, where the cancer has greater than 25 % expression level. CC modulators identified by (M1) are useful in a variety of diagnostic and therapeutic applications, where disease or disorder prognosis is related to defects in the p53 pathway, such as, angiogenesis, apoptotic or cell croliferation disorders (e.g. cancer). Another two new methods (M2 and CC also useful for treating defects in the p53 pathway of a cell, thus restoring the p53 function or progression through the cell cycle. (M2) and (M3) are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying modulators of the p53 pathway for use in treating apoptotic or cell proliferation disorders, comprises screening for agents that modulate activity of a human ortholog of genes that modify the p53 pathway in Drosophila.
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10-OCT-2001; 2001US-0328605F
15-FEB-2002; 2002US-0357253F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; p53 modifier; cytostatic; cancer; cytostatic; antiangiogenic; antiapoptotic; p53 pathway; breast cancer; colon cancer; kidney cancer; lung cancer; ovarian cancer; andjogenesis; cell cycle; apoptotic disorder; cell proliferation disorder.
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                        ACF12828 to ACF12947 encode the human cervical cancer marker proteins (I) CC given in ABR92047 to ABR92164. A higher level of expression of (I) than command indicates the presence of cervical cancer. Also described: (1) a cortaining (II) containing (II); (2) a host cell (III) containing (II); and (3) cassessing (MI) whether a patient is afflicted with cervical cancer. CC comprising comparing the level of expression of the marker in a patient's cc sample, and the normal level of expression of the marker in a control non cervical cancer sample, where a significant increase in the level of expression of the marker in the patient's sample relative to that in the control sample is an indication that the patient is afflicted with cervical cancer. (I) has cytostatic activity, and can be used in gene therapy and in vaccines. (I) is useful in detecting, characterising, correventing and treating human cervical cancers. (I) may also be used in various prognostic and diagnostic assays, pharmacogenomics and in control or control cancers are considered trials.
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Best Local
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                                                                                                                                                                                                                                                                                                                         New isolated nucleic acid molecule useful for detecting, characterizing, preventing and treating human cervical cancers, in various prognostic an diagnostic assays, in pharmacogenomics and in monitoring clinical trials
                                                                                                                                                                                                                                                                                            Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; ACF12941.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            apoptotic or cell proliferation disorders. The present sequence represents a human p53 pathway modifying protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-JUN-2001; 2001US-0298155P
13-JUN-2001; 2001US-0298159P
14-NOV-2001; 2001US-0335936P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-JUN-2002; 2002WO-US018638
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           detection; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human cervical cancer cell marker protein SEQ ID NO:226
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                                                                                                                                                                                                                                                                                            Page 377-378; 386pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Chen Y,
M, Glatt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer; cervical cancer marker; cancer therapy;
therapy; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Zhao X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monahan JE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14;
                                                                                                                                                                                                                                                                                                                             in monitoring clinical trials.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kamatkar
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Query Match Best Local Similarity

100.0%;

Score Pred.

No. DB 14;

6,

Length 105;

Sequence 105

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RESULT 60 ADD67269 ID ADD67

ADD67269; ADD67269 멍

31

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Matches

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Query Match
Best Local Similarity
6; Conserva
                                                                                                                                                                                                                                                           CC comprises: (a) capture agents, where each capture agent is preselected to comprising one of the pre-selected tag; and (b) tagged reagents, each comprising one of the pre-selected tag; where each locus in the comprises a molecule and a tag, each tag is pre-selected to specifically bind to a capture agent, where each tag is pre-selected to specifically comprises a molecule and a tag, each tag is pre-selected to specifically bind to a capture agent, where each tag is bound to a capture agent, comprises tagged reagent with the capture agent, where cach locus comprises tagged reagents and where each of the different complex of the tagged reagents and where each of the different complex at each locus comprises the same pre-selected tag. Also completes at each locus comprises the same pre-selected tag. Also completes at each locus comprises the same pre-selected tag. Also completes at each locus comprises the same pre-selected tag. Also completes (a) a method for profiling completes apples; (a) a method for profiling completes apples; (b) a method for profiling comprising comprising computer system that displays a collection of binding sites comprising capture agents bound to a solid support and tagged reagents; comprising capture agents bound to a solid support and tagged reagents; comprising a sample. The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New combination comprising an addressable collection of binding sites, software comprising instructions for pattern recognition and an imager for detecting patterns, useful for profiling a sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     apparatus combination; binding site collection; pattern recognition; profiling; screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprising: (a) an addressable collection of binding sites; and (b) software comprising instructions for pattern recognition and/or an imager for detecting patterns. The addressable collection of binding sites comprises; (a) capture agents, where each capture agent is preselected to
                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a combination of apparatus (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 7; SEQ ID NO 127; 309pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (POIN-) POINTILLISTE INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JAN-2003; 2003WO-US002397
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WCGPCK 6
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                                                                                                                                                                                                               105 AA;
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                                                                                         Conservative
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                                                                                                                        100.0%;
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                                                                                                                        Score 47; DB
Pred. No. 14;
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                                                                                                                                                 Length 105;
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ADF30728:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a method of treating an angiogenesis-mediated disorder in a subject. The method is useful for treating angiogenesis-mediated disorder, e.g., retinal or choroidal neovascularisation or diseases associated with chronic inflammation, myocardial ischaemia, stroke, coronary artery disease or peripheral vascular disease. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treating angiogenesis-mediated disorder, e.g., retinal or choroidal neovascularization or diseases associated with chronic inflammation, myocardial ischemia, stroke, coronary artery disease or peripheral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rat; angiogenesis; angiogenesis modulating protein; retinal neovascularisation; choroidal neovascularisation; chronic inflammation; myocardial ischaemia; stroke;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-FEB-2002; 2002US-0355295P
26-JUN-2002; 2002US-0391758P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 105
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N-PSDB; ADF30727.
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   mitochondrial; human; screening assay
Huntington's disease; osteoarthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 2; SEQ ID NO 291; 26pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vascular disease.
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                                                                                             Human heat mitochondrial protein as a therapeutic target SeqID2019.
                                                                                                                                                        06-MAY-2004
                                                                                                                                                                                                                                                                          ADJ70213 standard; protein; 105 AA.
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100.0%; Pr
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                               screening assay; diabetes mellitus;
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Pred. No.
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for therapeutic intervention in treating a disease associated with altered mitochondrial function. Specifically, it refers to a method for identifying protesins of the human heart mitochondrial proteome that are useful for drug screening assays, as well as therapeutic targets. The present invention describes a method for identifying such protesins that can be used in the treatment of various diseases associated with altered mitochondrial function including diabetes mellitus, Huntington's disease, osteoarthritis, Leber's hereditary optic neuropathy (LHON), mitochondrial encephalogathy lactic acidosis and stroke (MELAS), myoclonic epilepsy ragged red fibre syndrome (MERRF) or cancer. Accordingly, these compositions have neuroprotective, nootropic, antidiabetic, antiantiant and antiarthritic contrabile combination and and
Antidiabetic; Antiarteriosclerotic; Neuroprotective; Nootropic; Antiparkinsonian; Anticonvulsant; Cardiant; Cardiovascular; Antirheumatic; Antiarthritic; Cytostatic; Antipsoriatic; Gene T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          anticonvulsant, antiarthritic, osteopathic, ophthalmological and cytostatic activities. This polypeptide sequence is a human hear mitochondrial protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying a mitochondrial target for drug screening assays and treating diseases associated with altered mitochondrial function, comprises detecting a modified polypeptide in a sample and correlatives.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mitochondrial encephalopathy lactic acidosis and stroke; MELAS; myoclonic epilepsy ragged red fibre syndrome; MERR; cancer; neuroprotective; nootropic; antidiabetic; anticonvulsant; antiarthritic; osteopathic; ophthalmological; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ghosh SS,
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17-JUN-2002; 2002US-0389987P-
20-SEP-2002; 2002US-0412418P
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                                                                                                                                  06-MAY-2004
                                                                                                                                                                                                                  ADJ47593 standard; protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to novel mitochondrial targets that can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MITO-) MITOKOR.
(BUCK-) BUCK INST AGE RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Leber's hereditary optic neuropathy; LHON,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2003-845369/78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the disease
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                                                                                                                                                                                                                                                                                                                                                                       1 WCGPCK 6
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                                                                                     coding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2019; 180pp; English.
                                                                                                                               entry)
                                                                                     sequence, SEQ ID 1.
                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 47; Di
100.0%; Pred. No. 14
tive 0; Mismatches
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14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       7; Length 105;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 is a human heart
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Gene Therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          correlating
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  ADL82895
ID ADL828
XX ADL82
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Best Local (
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                                          Immunosuppressive;
Antiallergic; Musco
Gene Therapy; PRO;
                                                                                                                                                                          17-JUN-2004
                                                                                                                                                                                                                                                             ADL82895 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Min W,
                                                                                                                                  Human
                                                                                                                                                                                                                     ADL82895;
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                          lmmune-mediated
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                                                                                                                                                                                                                                                                                                                                                                       WCGPCK 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                          (first
                                          sive; Cytostatic; Antiarthritic; Antirheumatic; Antianemic;
Muscular; Neuroprotective; Nephrotropic; Antiinflammatory;
PRO; B cell related disorder; cancer;
                                                                                                                               SEQ
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protein;

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Gaps

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the oxidizing effects of cytokines or reactive oxygen species or resistant to S-nitrosylation of a SH-group by nitrous oxide. The mutant Trx proteins are useful for treating or diagnosing atherosclerosis, diabetes, apoptotic disease, mitochondrial dysfunction disease or cardiac dysfunction. Mitochondrial dysfunction diseases include Kearns-Sayre syndrome or encephalomyopathy lactic acidosis and stroke-like episodes. Apoptotic diseases include neurodegenerative diseases, e.g. Aliheimer's disease, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, multiple sclerosis or peripheral neuropathy. Cardiac dysfunction includes myocardial infarction, cardiomyopathy, arterial hypertension or heart failure. The mutant Trx proteins are also useful for treating or diagnosing angiogenesis dependent diseases, e.g. rheumatoid arthritis, cancer or psoriasis. The present sequence is the wild-type Trx protein, from which the mutants of the invention were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New mutant thioredoxin molecule which is resistant to oxidizing effects of cytokines or reactive oxygen species, useful for treating or diagnosing atheroselerosis, diabetes, Alzheimer's disease, myocardial infarction or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          thioredoxin; Trx; atherosclerosis; diabetes; apoptotic disease; mitochondrial disease; cardiac dysfunction; angiogenesis dependent disease; neurodegenerative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to mutant thioredoxin (Trx) proteins and their coding sequences, where the thioredoxin molecule is resistant to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 1; 145pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-157111/15.
N-PSDB; ADJ47596.
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                                    100.0%;
Score 47; DB Pred. No. 14; O; Mismatches
                                    DB
14;
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                                                                             Length 105
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RESULT 65
ADM32938
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             deficiency, selective deficiency of IgG subclasses, immunodeficiency with hyper IgM, transient hypogammaglobulinemia of infancy, Burkitt's lymphoma, intermediate lymphoma, follicular lymphoma, type II hypersensitivity, rheumatoid arthritis, autoimmune mediated haemolytic anaemia, myasthenia gravis, hypoadrenocorticism, glomerulonephritis, or ankylosing spondyllitis. The PRO proteins are also useful for preparing a medicament for treating a condition that is responsive to the PRO coding sequences are useful as hybridization probes in chromosome and gene mapping, in preparing PRO proteins, or in generating transgenic animals or knockout animals, which in turn are useful in the development and screening of therapeutically useful reagents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New PRO polypeptide, useful for diagnosing and treating a B cell related disorder, e.g. Burkitt's lymphoma, rheumatoid arthritis, autoimmune mediated hemolytic anemia, myasthenia gravis or ankylosing spondylitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 10; Fig 97; 695pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chiu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-SEP-2002; 2002US-0411392P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-SEP-2003; 2003WO-US029097
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                                                                                                                                         mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive sputum; lung disease; cystic fibrosis; enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antigen unresponsiveness, selective IgA deficiency, selective IgM deficiency, selective deficiency of IgG subclasses, immunodeficiency with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to PRO proteins and their coding sequences. The PRO proteins are useful for diagnosing and treating a B cell related disorder, e.g. X-linked infantile hypogammaglobulinemia, polyeaccharide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wu TD;
10-SEP-2003; 2003WO-US028526.
                               25-MAR-2004
                                                              WO2004024868-A2
                                                                                                          Homo sapiens
                                                                                                                                                                                                                      17-JUN-2004
                                                                                                                                                                                                                                                                                   ADM32938 standard; protein; 105 AA.
                                                                                                                                                                                         Amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GETH )
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                                                                                                                                                                                                                                                                                                                                                                                               1 WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Clark H, Dennis K,
                                                                                                                                                                                       sequence of a thioredoxin polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 47; DB
100.0%; Pred. No. 14;
                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fong S,
                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                            0;
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                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                              0,
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mucus; sputum; thioredoxin; active site; cohesive lung disease; cystic fibrosis; enzyme. Amino acid sequence of a thioredoxin polypeptide.

mucus; cohesive sputum;

Mus musculus. Synthetic.

25-MAR-2004

WO2004024868-A2

10-SEP-2003; 2003WO-US028526

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Conservative

0;

Indels

<u>.</u>

Gaps

0

ADM32935 RESULT 66

ADM32935 standard; protein; 105

17-JUN-2004 ADM32935;

(first entry)

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The specification describes a method for increasing the liquefaction of CC mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum. The method comprises contacting the mucus or sputum of the patient with a composition comprises contacting the mucus or sputum of contacted is located in the respiratory tract, gastrointestinal tract or contacted is located in the respiratory tract, gastrointestinal tract or patient in a pharmaceutical carrier. The composition is administered to the contacted in a pharmaceutical carrier. The protein has a half-life in the contacted contacted in the respiratory tract, gastrointestinal tract or the contacted in a pharmaceutical carrier. The protein has a half-life in the contacted of a sample of mucus or sputum from the patient shows a contacted from the patient of a sample of mucus or sputum from the patient shows a contacted contacted in the protein comprises thioredoxin and mammalian in the composition. The protein comprises after administration of the composition. The composition further comprises nicotinamide-adenine contacted from phosphate (reduced from) (NADPH) for reducing the contacted phosphate (reduced from) (NADPH) for reducing the contacted phosphate (reduced form) in a pharmaceution of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and contestive viscosity of contestive mucus or sputum. These may be used for contestive viscosity of contestiveness of mucus or sputum is a sumptom or contestive respective respective respectiveness of mucus or sputum is a sumptom or contestive respective respectiveness of mucus or sputum.
Query Match
Best Local Similarity
Matches 6; Conser
                                                                                                                                                            excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. ADM32930ADM32941 represent thioredoxin polypeptiwhich may used in the method of the invention.
                                                                                                       Sequence 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 66-67; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprising a protein or peptide containing a thioredoxin active-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Increasing the liquefaction of excessively viscous or cohesive mucus sputum comprises contacting the mucus or sputum with a composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-270016/25.
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11-APR-2003; 2003US-0462082P.
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100.0%; Score 47; DI
100.0%; Pred. No. 14
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                                                     Length 105;
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RESULT 67
ADM32936
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total volume of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian thioredoxin. The composition further comprises nicotinamide-adenine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 dinucleotide phosphate (reduced form) (NADPH) for reducing the thioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for treating lung diseases (i.e. cystic fibrosis) in which abnormal or excessive viscosity or cohesiveness of mucus or sputum is a symptom or cause of the disease. ADM32930ADW32941 represent thioredoxin polypeptides which may used in the method of the invention.
                                                                                                                                                               mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive lung disease; cystic fibrosis; enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The specification describes a method for increasing the liquefaction of mucus or sputum in a patient that has excessively viscous or cohesive mucus or sputum. The method comprises contacting the mucus or sputum of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Increasing the liquefaction of excessively viscous or cohesive mucus or sputum comprises contacting the mucus or sputum with a composition comprising a protein or peptide containing a thioredoxin active-site in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-SEP-2002; 2002US-0409960P.
11-APR-2003; 2003US-0462082P.
10-SEP-2003; 2003WO-US028526
                                                                                                                            Rattus norvegicus.
                                                                                                                                                                                                                                                          17-JUN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                  ADM32936 standard; protein; 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 65; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-270016/25
                                                                       WO2004024868-A2
                                                                                                                                                                                                                     Amino acid sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                     of a thioredoxin polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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Pred. No. 14;
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8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 105;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                    sputum;
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RX 11-SEP-2002; 2002US-0403960P.
RR 11-APR-2003; 2003US-0462082P.
XX (NAJE-) NAT JEWISH MEDICAL & RES CENT.
XX (NAJE-) NAT JEWISH MEDICAL & RES CENT.
XX (WPI; 2004-270016/25.
XX Fincreasing the liquefaction of excessively viscous or cohesive mucus or sputum comprises contacting the mucus or sputum with a composition comprises contacting the mucus or sputum with a composition of reduced state.
XX Disclosure; Page 65-66; 69pp; English.
XX Commous or sputum. The method comprises contacting the mucus or sputum of mucus or sputum. The method comprises contacting the mucus or sputum of mucus or sputum of mucus or sputum. The method comprises contacting the mucus or sputum of mucus or sputum of thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administration of the patient in a pharmaceutical carrier. The procein has a half-life in the patient of about 5 minutes to about 24 hours. The liquid phase of a total volume of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the patient in phase of a total composition. The procein has a half-life in the composition and located from prokaryotic thioredoxin, past thioredoxin, plant thioredoxin selected from prokaryotic chioredoxin, plant plant thioredoxin medicade. The composition are useful for decreasing the viscosity and cohesive mucus or sputum. These may be used for a sample of the disease. AMM32930AMM3201 represent thioredoxin
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                                 Matches
                                        Query Match
Best Local Similarity
31
                1 WCGPCK 6
                                 6,
WCGPCK 36
                                 Conservative
                                         100.0%; Score 47; DB 100.0%; Pred. No. 14;
                                 ..
                                 Mismatches
                                                 BB
                                                 8;
                                 <u>,</u>
                                                Length 105,
                                 0;
                                 Gaps
                                 0
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RESULT 68
ADM32937
ID ADM32937 standard; protein; 105 AA.

XX
AC ADM32937;
XX
DT 17-JUN-2004 (first entry)
XX
DE Amino acid sequence of a thioredoxin polypeptide.
XX
XX
XX
XX
XX
XX
XX
XX
XX
Iung disease; cystic fibrosis; enzyme.
XX
OS Bos taurus.
OS Synthetic.
XX
PN W02004024868-A2.
XX
PP 25-MAR-2004.
XX
PF 10-SEP-2003; 2003WO-US028526.
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ADM32934
ID ADM1
XX
AC ADM1
XX
DT 17-1
XX
DT 17-1
XX
DT 17-1
XX
IN MUCI
KW MUCI
KW MUCI
KW Iuni
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Syn
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PN WO2
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PF 10-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The specification describes a method for increasing the liquefaction of CC mucus or sputum in a patient that has excessively viscous or cohesive CC mucus or sputum. The method comprises contacting the mucus or sputum of CC the patient with a composition comprising a protein or peptide containing CC a thioredoxin active site in reduced state. The mucus or sputum to be CC contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the CC patient in a pharmaceutical carrier. The protein has a half-life in the CC patient of about 5 minutes to about 24 hours. The liquid phase of a total CC volume of a sample of mucus or sputum from the patient shows a CC statistically significant increase after administration of the CC composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian CC intoredoxin. The composition further comprises nicotinamide-adenine CC dinucleotide phosphate (reduced form) (NADPH) for reducing the thioredoxin reductase. The CC method and composition are useful for decreasing the viscosity and composition are useful for decreasing the viscosity and composition are useful for decreasing the viscosity or abnormally viscous or cohesive mucus or sputum. These may be used for treating lung diseases (i.e. cystic fibrosis) in which abnormal or content of the protein or sputum.
                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 69
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Best Local &
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11-APR-2003; 2003US-0462082P.
  10-SEP-2003; 2003WO-US028526
                                          25-MAR-2004
                                                                                WO2004024868-A2
                                                                                                                                          Gallus
                                                                                                                                                                                mucus; sputum; thioredoxin; active site; cohesive mucus; cohesive lung disease; cystic fibrosis; enzyme.
                                                                                                                                                                                                                                              Amino acid sequence of a thioredoxin polypeptide.
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Best Local Matches

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Human PRO polypeptide #292.

12-AUG-2004

(first entry)

Human; PRO; immune related disorder; systemic lupus erythematosus; rheumatoid arthritis; osteoarthritis; juvenile chronic arthritis; systemic scherosis; Sjogren's syndrome; vasculitis; sarcoidosis; autoimmune haemolytic anaemia; autoimmune thrombocytopenia; thyroiditis; diabetes mellitus; renal disease; demyelinating disease; central nervous system; peripheral nervous system; demyelinating polyneuropathy; Guillain-Barre syndrome;

demyelinating

polyneuropathy.

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The specification describes a method for increasing the liquefaction of commucus or sputum. The method comprises contacting the mucus or sputum of the patient with a composition comprises contacting the mucus or sputum of the patient with a composition comprising a protein or peptide containing a thioredoxin active site in reduced state. The mucus or sputum to be contacted is located in the respiratory tract, gastrointestinal tract or reproductive tract of the patient. The composition is administered to the patient in a pharmaceutical carrier. The protein has a half-life in the composition of a sample of mucus or sputum from the patient shows a statistically significant increase after administration of the composition. The protein comprises thioredoxin selected from prokaryotic thioredoxin, yeast thioredoxin, plant thioredoxin and mammalian thioredoxin. The composition further comprises nicotinamide-adenine thioredoxin active site of the protein, and thioredoxin reductase. The chioredoxin active site of the protein, and thioredoxin reductase. The method and composition are useful for decreasing the viscosity and cohesiveness of, and increasing the liquefaction of excessively or abnormally viscous or cohesive mucus or sputum. These may be used for treating lung diseases (i.e. cystic fibrosis) in which abnormal or cause of the disease. ADM32930ADM32941 represent thioredoxin polypeptides which may used in the method of the invention.
                                             Query Match
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                                                                                                                                                                                                                                                                      The invention relates to human PRO polypeptides and the polynucleotides encoding them. The polypeptides and polynucleotides are useful for treating and diagnosing immune related disorders in mammals. The immune related disorders include systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, systemic sclerosis, Sjogren's syndrome, vasculitis, sarcoidosis, autoimmune haemolytic anaemia, autoimmune thrombocytopenia, thyroiditis, diabetes mellitus, immune-mediated renal disease, demyelinating diseases of the central or peripheral nervous system, demyelinating polyneuropathy, Guillain-Barre syndrome and chronic inflammatory demyelinating polypeptide of the invertice.
                                                                                                                                                                                                                        Sequence 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel PRO polypeptide e.g., PRO69614, PRO71106, or PRO86388 useful for treating an immune related disorder such as systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis or spondyloarthropathy.
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N-PSDB; ADO19654.
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A;Authors: Hou, S.; Daniels, C.J.; Dennis, P.P.; Omer, A.D.; Ebhardt, H.; Lowe, T.M.; A;Title: Genome sequence of Halobacterium species NRC-1.
A;Reference number: A94160; MUID:20504483; PMID:11016950
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Best Local Similarity
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A,Residues: 1-102 <TIGR>
A,Cross-references: UNIPROT:P47370; UNIPARC:UPI0000136E89; GB:U39691; GB:L43967; NID:g10
A,Experimental source: strain G-37
C;Genetics:
A,Genetic code: SGC3
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: redox-active disulfide
C;Keywords: redox-active disulfide
C;Keywords: thioredoxin homology <THR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: Complete sequence analysis of the genome of the bacterium Mycoplasma A;Reference number: $73327; MUID:97105885; PMID:8948633

A;Accession: $73896

A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: DNA

A;Residues: 1-102 <HIM>
A;Cross-references: UNIPROT:P75512; UNIPARC:UPI0000136E8A; EMBL:AE000056; GB:UA;Cross-references: UNIPROT:P75512; UNIPARC:UPI0000136E8A; EMBL:AE000056; GB:UA;Note: the nucleotide sequence was submitted to the EMBL Data Library, NovembC;Genetics:
                                                                                                               A;Status: preliminary
A;Molecule type: DNA
A;Moseidues: 1-102 <ARN>
A;Cross-references: UNIPROT:084544; UNIPARC:UPI0000136E75; GB:AE001324; GB:AE001273; NII
A;Experimental source: serotype D, strain UW-3/Cx
C;Genetics:
                                                                                                                                                                                                                                                                                                                    probable thioredoxin - Chlamydia trachomatis (serotype D, strain UW3/Cx) C;Species: Chlamydia trachomatis C;Date: 13-Sep-1998 #sequence_revision 13-Sep-1998 #text_change 05-Oct-2 C;Accession: B71503  
R;Stephens, R.S.; Kalman, S.; Lammel, C.J.; Fan, J.; Marathe, R.; Aravin Science 282, 754-759, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 4
B71503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N;Alternate names: hypothetical protein A65_orf102
C;Species: Mycoplasma pneumoniae
A;Variety: ATCC 29342
C;Date: 27-Feb-1997 #sequence_revision 25-Apr-1997 #text_change 09-Jul-2004
C;Accession: S73896
                                                                                                                                                                                                                                                    A;Title: Genome sequence of A;Reference number: A71570; A;Accession: B71503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Genetic code: SGC3
C; Superfamily: thior
F; 9-90/Domain: thior
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Himmelreich, R.; Hilbert, H.; Plagens, H.; Pirkl, Nucleic Acids Res. 24, 4420-4449, 1996
                                                                    A;Gene: trxA
C;Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           F;30-33/Disulfide bonds: redox-active #status predicted
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    Mycoplasma pneumoniae (strain ATCC 29342)
    names: hypothetical protein A65_orf102

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                                                                       thioredoxin; thioredoxin homology
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                                                                                                                                                                                                                                                                                of an obligate intracellular pathogen 70; MUID:99000809; PMID:9784136
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Pred. No.
    Score
Pred.
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                          Length 102
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R;Muller, E.G.D.
J. Biol. Chem. 266, 9194-9202, 1991
A;Title: Thioredoxin deficiency in yeast prolongs S phase and A;Reference number: A39847; MUID:91225027; PMID:2026619
A;Accession: A39847
                                                              A;Cross-references: UNIPARC:UPI0000171E21 R;Koetter, P.; Rose, M.; Entian, K.D. submitted to the Protein Sequence Database,
                                                                                                                                                                   A;Reference number: S05793; MUID:72100583; PMID:4945270
A;Accession: S15360
                                                                                                                                                                                                                                             A;Cross-references: UNIPARC:UPI000004F943; R;Hall, D.E.; Baldesten, A.; Holmgren, A.; Eur. J. Biochem. 23, 328-335, 1971
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                   A; Reference number: S64863
A; Accession: S64870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Title: Yeast thioredoxin genes.
A; Reference number: A38669; MUID:91107668; PMID:1988444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thioredoxin II - yeast (Saccharomyces cerevisiae)
N;Alternate names: protein L1933; protein YLR043c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPROT:Q9PJK3; UNIPARC:UPI0000057A90; GB:AE002349; GB:AE002160; NII
A;Experimental source: strain Nigg (MoPn)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Reference number: A81500; MUID:20150255; PMID:10684935
A;Accession: C81660
                                                                                                                                                       A; Molecule type: protein
                                                                                                                                                                                                                                                                                                                          A; Residues: 1-103 <MUL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-103 <GAN>
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A; Residues: 1-102 <TET>
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                                                                                                                               A; Residues: 26-34 <HAL>
                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;Species: Saccharomyces cerevisiae
;Date: 31-Dec-1991 #sequence_revision 31-Dec-1991 #text_change 05-Oct-2004
;Accession: S15048; A39847; S15360; S64870; B38669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hioredoxin TC0826 [imported] - Chlamydia muridarum (strain Nigg)
;Species: Chlamydia muridarum, Chlamydia trachomatis MoPn
;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 05-Oct-2004
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                                                                May 1996
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Salzberg,
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A; Reference number: A; Accession: A28086
                   R;Johnson, R.S.; Mathews, W.R.; Biemann, K.; Hopper, S. J. Biol. Chem. 263, 9589-9597, 1988
A;Title: Amino acid sequence of thioredoxin isolated from A;Reference number: A28086; MUID:88257078; PMID:3164311
                                                                                                                                 C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 30-Jun-1989 #sequence_revision 26-May-1994
                                                                                                                                                                                                                               RESULT 8
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                                                                                                              C;Accession: A28086
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thioredoxin II - fission yeast (Schizosaccharomyces pombe)
N,Alternate names: thioredoxine 2
C;Species: Schizosaccharomyces pombe
C;Date: 03-Dec-1999 #sequence revision 03-Dec-1999 #text_change 09-Jul-2004
C;Accession: T39085; T46567
                                                                                                                                                                                                                                                                                                         A; Map position: 1
C; Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPARC:UPI000016205F; EMBL:AJ003819; A;Experimental source: strain 972(-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Description: TRX2, a fission yeast stress protein. A;Reference number: Z23075 A;Accession: T46567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Experimental source: strain 972h-; cosmid c7D4 R;Lenaers, G.; Perret, E.; Bonin, O.; Picard, A.; Caput, submitted to the EMBL Data Library, October 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Reference number: Z21826
A; Accession: T39085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          F;2-103/Product: thioredoxin II #status predicted <MAT>F;9-90/Domain: thioredoxin homology <THR>F;30-33/Disulfide bonds: redox-active #status experimen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Map position: 12R C; Superfamily: thioredoxin; thioredoxin homology
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A;Cross-references: SGD:S0004033; MIPS:YLR043c
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A;Cross-references: UNIPARC:UPI000004F943; EMBL:Z73215; NID:g1360372; PIDN:CAA97572.1; |
A;Note: experimental_source strain S288C
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                              A;Gene: trx2; SPAC7D4.07c
                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Residues: 1-103 < LEN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Status: preliminary; translated from GB/EMBL/DDB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Keywords: redox-active disulfide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
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rabbit
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Pred. No.
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                                                                                                                                                                                                                   Mismatches
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#text_change 05-Oct-2004

rabbit bone marrow determined

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A; Residues: 1-104 C/HE>
A; Cross-references: UNIPROT: P14949; UNIPARC: UPI00006087B; GB: J03294; GB: M26384; NID: g14
A; Cross-references: UNIPROT: P14949; UNIPARC: UPI00006087B; GB: J03294; GB: M26384; NID: g14
A; Cross-references: UNIPROT: P14949; UNIPARC: UPI00006087B; GB: J03294; GB: M26384; NID: g14
A; Cross-references: UNIPROT: P14949; UNIPARC: UPI00006087B; GB: J03294; GB: M26384; NID: g14
A; Cross-references: University C.; Fullian, K.D.; Errington, J.; Fazevedo, V.; M.; Chc
A; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A; Authors: Dulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Galler
iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F.
Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois,
A; Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maueel
Y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; Portetelle
Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Scanlon,
A; Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, A.; Seror
akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terspatra, P.; Tognoni, A.; Tosato, V.; Uchiyama,
T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yoshida, K
A; Authors: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A.
A; Title: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis.
A; Accession: H69726
A; Catalega, C.; Folia, C.; Foli
                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Map position: 70 min
C;Superfamily: thioredox
C;Keywords: redox-active
F;9-90/Domain: thioredox
F;29-32/Disulfide bonds
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C;Date: 31-Uan-1992 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: B37192; H69726
R;Chen, N.Y; Zhang, J.J.; Paulus, H.
J. Gen. Microbiol. 135, 2931-2940, 1989
A;Title: Chromosomal location of the Bacillus subtilis aspartokinase II general A;Reference number: A37192; MUID:90132525; PMID:2559145
A;Accession: B37192
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thioredoxin
                                       TXBY1
                                                                    RESULT 10
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A; Cross-references: UNII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
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Best Local
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in: thioredoxin homology <THR>
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yeast
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   cerevisiae)
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   A; Map position: 7R
C; Superfamily: thioredoxin; thioredoxin homology
C; Keywords: redox-active disulfide
                                                                                                     A;Gene: SGD:TRX2; TR-I; MIPS:YGR209c
A;Cross-references: SGD:S0003441; MI
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es, of the human:

A;Reference number: $53922

A;Accession: $53922

A;Accession: $53932

A;Accession: $53932

A;Accession: $53932

A;Accession: $53932

A;Accession: $53932

A;Accession: $53932

A;Coss-references: UNIPARC:UPIO00004F91C; EMBL:Z49133; NID:g790489; PIDN:CAA89002.1; PJA;Cross-references: UNIPARC:UPIO00004F91C; EMBL:Z49133; NID:g790489; PIDN:CAA89002.1; PJA;Cross-references: UNIPARC:UPIO00004F91C; EMBL:Z49133; NID:g790489; PIDN:CAA89002.1; PJA;Accession: Analysis of the 15.6-kb fragment encompassing the ADE3 gene.

A;Reference number: $61947

A;Reference number: $61947

A;Reference number: $61947

A;Reference number: $61947

A;Residues: 1-104 <SON>
A;Residues: 1-104 <SON>
A;Residues: 1-104 <SON>
A;Residues: 1-104 <SON>
A;Cross-references: UNIPARC:UPIO00004F91C; EMBL:U40843; NID:g1165213; PIDN:AAA85584.1; LA;Cross-references: UNIPARC:UPIO00004F91C; EMBL:U40843; NID:g1165213; PIDN:AAA885584.1; LA;Cross-references: UNIPARC:UPIO00004F91C; EMBL:U40843; NID:g1165213; PID
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A;Description: Sequencing of a 17.6 kb segment on the right arm of yeast chromosome VII
                                                                                                                                                                                                                                                                                                                                                                                 Yeast 12, 273-280, 1996
Yeast 12, 273-280, 1996
A;Title: Sequencing of a 17.6 kb segment on the right arm of yeast terial electron-transferring flavoproteins (beta-chain) and of the terial electron transferring flavor flavor
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A;Residues: 1-104 <GUW>
A;Residues: 1-104 <GUW>
A;Residues: 1-104 <GUW>
A;Residues: 1-104 <GUWA
A;Experimental source: strain S288C
A;Experimental source: strain S288C
R;Guerreiro, P.; Barreiros, T.; Soares, H.; Cyrne, L.; Maia e Silva, A.; Rodrigues-Pous
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A;Title: Yeast thioredoxin genes.
A;Reference number: A38669; MUID:91107668;
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C;Date: 31-Dec-1991 #eequence revision 31-Dec-1991 #text change 05-Oct-2004
C;Accession: S15049; B39847; S05793; S53932; S61947; S64531; S63858; A38669
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                                                                  A; Molecule type: DNA
A; Residues: 1-104 < GUF >
A; Cross-references: UNI
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A; Accession: S64531
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A; Residues: 1-104 < GAN>
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       A; Note: the nucleotide
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                                                                                                                                                                                                                                                                                        acid sequence not shown; translation not shown
                                                                         UNIPARC: UPI000004F91C; EMBL: Z49133; NID: G790489; PIDN: CAA89002.1;
sequence
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       submitted to the EMBL Data Library, April 1995
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A;Cross-references: UNIPARC:UP10000174DEF
R;Hammel, K.E.; Cornwell, K.L.; Buchanan, B.B.
Proc. Natl. Acad. Sci. USA 80, 3681-3685, 1983
A;Title: Ferredoxin/flavoprotein-linked pathway for the reduction of thioredoxin.
                                                                                                                                        A;Cross-references: UNIPARC:UPI0000174DEF
R;Reynolds, C.M.; Poole, L.B.; Hammel, K.E.;
submitted to the Protein Sequence Database, S
A;Reference number: A59394
A;Accession: A59394
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R;Bork, P.; Ouzounis, C.; Casari, G.; Schneider, R.; Sander, C.

Mol. Microbiol. 16, 955-967, 1995

A;Title: Exploring the Mycoplasma capricolum genome: a minimal A;Reference number: S77739; MUID:96059641; PMID:7476192

A;Accession: S77780
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N;Alternate names: protein MC064
                                                                                A; Molecule type: protein A; Residues: 1-104 < RE2 >
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A;Accession: B59394
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                                                                                                                        A;Status: preliminary
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13-93/Domain: thioredoxin homology <THR>
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100.0%; Pred. No. 1;
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C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K. Lancet 357, 1225-1240, 2001
A;Title: Whole genome sequencing of meticillin-resistant Stapylococcus aureus. A;Reference number: A89758; MUID:21311952; PMID:11418146
A;Accession: E89885
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                                                                                                                                                                              C; Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                            C; Genetics:
                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:Q9ZEH4; UNIPARC:UPI00000D78B6; GB:BA000018; PID:g13700945; A;Experimental source: strain N315
                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-104 < KUR>
                                                                                                                                                                                                                                                                                                                                            A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 05-Oct-2004 C;Accession: E89885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; Species: Staphylococcus aureus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 thioredoxin [imported] - Staphylococcus aureus (strain N315)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Experimental source: strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-104 < STO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Complete genome sequence of the alkaliphilic bacterium Bacillus halodurans A;Reference number: A83650; MUID:20512582; PMID:11058132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Takami, H.; Nakasone, K.; Takaki, Y.; Maeno, Nucleic Acids Res. 28, 4317-4331, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       thioredoxin trxA [imported] - Bacillus halodurans (strain C-125)
C;Species: Bacillus halodurans
C;Date: 01-Dec-2000 #sequence_revision 01-Dec-2000 #text_change 05-Oct-2004
C;Accession: B84037
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                                                                                        Matches
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                                                                                                                                   Query Match
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R; Martin, H.; Dean, M. Biochem. Biochem. Biochem. Commun. 175, 123-128, 1991
A; Title: Identification of a thioredoxin-related protein associated with plasma A; Reference number: PT0079; MUID:91151337; PMID:1998498
A; Accession: PT0079
                                                                                                A;Cross-references: UNIPARC:UPI000011065C; EMBL:X77584; NID:g453963; PIDN:CAA54687.1; PIR:Wollman, E.E.; d'Auriol, L.; Rimsky, L.; Shaw, A.; Jacquot, J.P.; Wingfield, P.; Grabe J. Biol. Chem. 263, 15506-15512, 1988
A;Title: Cloning and expression of a cDNA for human thioredoxin.
A;Reference number: A31993; MUID:89008454; PMID:3170595
A;Accession: A31993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Homo sapiens (man)
C;Date: 30-Jun-1992 #sequence revision 26-May-1994 #text change 05-Oct-2004
C;Accession: JH0568; S04106; S44375; A31993; PT0079; A60749; A38922; S53453; R;Tonissen, K.F.; Wells, J.R.B.
Gene 102, 221-228, 1991
A;Title: Isolation and characterization of human thioredoxin-encoding genes. A;Reference number: JH0568; MUID:91340156; PMID:1874447
A;Accession: JH0568
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J. Biol. Chem. 263, 9607-9611, 1988
A;Title: Isolation of a chicken thioredoxin cDNA clone:
A;Reference number: A30006; MUID:88257080; PMID:2838473
A;Accession: A30006
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A; Residues: 1-105 < JON>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Title: ATL-derived factor (ADF), an IL-2 receptor/Tac A;Reference number: S04106; MUID:89251607; PMID:2785919 A;Accession: S04106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: UNIPROT:P10599; UNIPARC:UPI000011065C; EMBL:X54539; NID:g37455; PIDN R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kondo, N.; Matsui, H.; Hamuro, J.; Brown, N.; Arai EMBO J. 8, 757-764, 1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: DNA
A;Residues: 1-105 <TON>
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;Date: 31-Mar-1989 #sequ
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Residues: 1-105 <TAG2>
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100.0%; Pred. No.
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                                                                                                                                                  GB:J04026; NID:g339648; PIDN:AAA74596.1;
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A; Molecule type: protein

A; Residues: 1-21;38-57 < DEA>

A; Residues: 1-21;38-57 < DEA>

A; Residues: 1-21;38-57 < DEA>

A; Cross-references: UNIPARC: UPI00000353C5; UNIPARC: UPI0000171E1C

A; Note: described to be a surface-associated thioredoxin

R; Wakasugi, H.; Rimsky, L.; Mahe, Y.; Kamel, A.M.; Fradelizi, D.; Tur.

Proc. Natl. Acad. Sci. U.S.A. 84, 804-808, 1987

A; Title: Epstein-Barr virus-containing B-cell line produces an interl.

A; Reference number: A60870; MUID: 87118252; PMID: 3027706

A; Contents: annotation

R; Weichsel, A.; Gasdaska, J.R.; Powis, G.; Montfort, W.R.

submitted to the Brookhaven Protein Data Bank, February 1996

A; Contents: annotation; X-ray crystallography, 1.7 angstroms, reduced

R; Weichsel, A.; Gasdaska, J.R.; Powis, G.; Montfort, W.R.

Schmitted, A.; Gasdaska, J.R.; Powis, G.; Montfort, W.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     submitted to the Brookhaven Protein Data Bank, February 1996
A;Reference number: A65534; PDB:1ERU
A;Contents: annotation; X-ray crystallography, 2.1 angstroms, oxidized forn
R;Forman-Kay, J.D.; Clore, G.M.; Gronenborn, A.M.
submitted to the Brookhaven Protein Data Bank, December 1990
A;Reference number: A50924; PDB:4TRX
A;Rontents: annotation; conformation by (1)H-NNR, residues 1-73,'T',75-105
R;Forman-Kay, J.D.; Clore, G.M.; Wingfield, P.T.; Gronenborn, A.M.
Biochemistry 30, 2685-2698, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Dean, M.F.; Martin, H.; Sansom, P.A.
Biochem. J. 304, 861-867, 1994
A;Title: Characterization of a thioredoxin-related surface protein.
A;Reference number: S53453; MUID:95118305; PMID:7818492
A;Accession: S53453
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A;Residues: 2-12, 'K',14-15, 'XX',18-19, 'X',21-22 <SIL>
A;Cross-references: UNIPARC:UPI0000171E1A
A;Cross-references: UNIPARC:UPI0000171E1A
A;Note: the abstract is inconsistent with figure 4 in having one undetermined residue af R;Rimsky, L.; Wakasugi, H.; Ferrara, P.; Robin, P.; Capdevielle, J.; Tursz, T.; Fradeliz J. Immunol. 136, 3304-3310, 1986
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R;Silberstein, D.S.; Ali, M.H.; Baker, S.L.; David, J.R.
J. Immunol. 143, 979-983, 1989
A;Title: Human eosinophil cytotoxicity-enhancing factor.
A;Reference number: A60749; MUID:89309777; PMID:2745979
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                                                                                                                                                                                                                                                                                                                                                    A;Map position: 9q31-9q31
A;Introns: 8/3; 43/3; 63/3; 85/3
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: High-resolution three-dimensional structure of reduced recombinant A;Reference number: A38953; MUID:91159399; PMID:2001356 A;Contents: annotation; conformation by (1)H- and (15)N-NMR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: UNIPARC:UPI0000171E1B R;Dean, M.F.; Martin, H.; Sansom, P.A.
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                                                                                                                                                                                                                                                                                      C;Keywords: redox-active disulfide F;2-105/Product: thioredoxin #status experimental
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   1 WCGPCK 6
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                                                                                                          Pred. No.
                                                                                                                                           Score 47;
                                                                            Mismatches
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A; Cross-reference.

A; Cross-reference.

A; Cross-reference.

A; Map position: 4:24.6

A; Introns: 29/2; 44/1; 84/2

C; Superfamily: thioredoxin; thioredoxin homology

C; Keywords: redox-active disulfide

C; Keywords: redox-active disulfide

C; Keywords: redox-active #status predicted

A7: DB 1; Le
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A;Title: Structure of the mouse thioredoxin-encoding gene and its processed pseudogene.
A;Reference number: JC4068; MUID:95137382; PMID:7835695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Mus musculus (house mouse)
C;Date: 21-Nov-1993 #sequence_revision 17-Oct-1997 #text_change
C;Accession: JC4068; S44376; S04107
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A;Title: Thioredoxin gene expression is transcriptionally up-regulated by retinol A;Reference number: JS0667; MUID:92181438; PMID:1543487
A;Accession: JS0667
                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: mRNA
A;Residues: 1-93,'N',94-96,'ALT',100-104,'S' <TAG2>
A;Cross-references: UNIPARC:UPI0000171E1E; GB:X77585
                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Title: ATL-derived factor (ADF), an IL-2 receptor/Tac A;Reference number: S04106; MUID:89251607; PMID:2785919 A;Accession: S04107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: UNIPARC:UPI0000163891; EMBL:X77585; NID:g453971; R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kondo, N.; Matsui, H.; Hamuro, EMBO J. 8, 757-764, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPROT:P10639; UNIPARC:UP10000163891; DDBJ:D21855; R;Tagaya, Y.; Maeda, Y.; Mitsui, A.; Kando, N.; Matsui, H.; Hamuro, J.; EMBO J. 13, 2244, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  thioredoxin - mouse
N;Alternate names: ATL-derived factor (ADF)
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A; Residues: 1-105 < ANG>
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C;Species: Macaca mulatta (rhesus macaque)
C;Date: 30-Jun-1992 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
C;Accession: JS0667
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A; Residues: 1-105 < TAG1>
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                    Similarity 6; Conserv
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                  100.0%; Score 47; llarity 100.0%; Pred. No. Conservative 0; Mismatch
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Brown, N.;
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RESULT 19
S04352
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A;Accession: A00281
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c;Species: coryneform bacterium ATCC11425
C;Date: 02-Apr-1982 #sequence_revision 02-Apr-1982 #text_change 05-Oct-2004
C;Accession: A00281
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Nucleic Acids Res. 17, 3973, 1989
A;Tille: Nucleotide sequence of a cDNA encoding rat thioredoxin.
A;Reference number: S04352; MUID:89282399; PMID:2734107
                                                                                                                                                            C;Keywords: redox-active disulfide F;8-91/Domain: thioredoxin homology <THR>
                                                                                                                                                                                                        A;Cross-references: UNIPROT: P00275; UNIPARC: UPT10000136D69
A;Note: the source was designated as Corynebacterium nephridii C;Comment: Thioredoxins are ubiquitous small hydrogen carrier C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                           A; Molecule type: protein A; Residues: 1-105 < MEN>
                                                                                                                                                                                                                                                                                                                                                                                                       R;Meng, M.; Hogenkamp, H.P.C.
J. Biol. Chem. 256, 9174-9182, 1981
A;Title: Purification, characteriza
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Biochem. J. 304, 861-867, 1994
A;Title: Characterization of a thioredoxin-related surface protein.
A;Reference number: S53453; MUID:95118305; PMID:7818492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:P11232; UNIPARC:UPI000003066A; EMBL:X14878; NID:g57385; PIDNR;Dean, M.F.; Martin, H.; Sansom, P.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-105 < TON>
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Pred. No. 1;
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WCGPCK 34

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cnloredoxin [imported] - Clostridium acetobutylicum
C;Species: Clostridium acetobutylicum
C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 05-Oct-2004
C;Accession: D97279
R;Nolling, J.; Breton, G.: Omelohanko """
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A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-105 <KUR>
A;Cross-references: UNIPROT:Q97EM7; UNIPARC:UPI00000CA715; GB:AE001437; PIDN:AAK81023.1
A;Cross-references: UNIPROT:Q97EM7; UNIPARC:UPI00000CA715; GB:AE001437; PIDN:AAK81023.1
A;Cross-references: UNIPROT:Q97EM7; UNIPARC:UPI00000CA715; GB:AE001437; PIDN:AAK81023.1
                                                                                                                                                                                                                                                          RESULT 23
S33357
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A;Title: Mechanisms of Evolution in Rickettsia conorii and Rickettsia prowazekii.
A;Reference number: A97700; MUID:21442074; PMID:11557893
A;Accession: B97700
                                                                                                                   thioredoxin - Streptomyces aureofaciens (fragment) C;Species: Streptomyces aureofaciens C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change C;Accession: S3357 C;Accession: S3357 R;Labudova, O; Nemethova, M.; Turna, J.; Kollarova, M. submitted to the EMBL Data Library, April 1993
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C;Species: R
A;Reference number: S33357
A;Accession: S33357
A;Molecule type: DNA
A;Residues: 1-106 <LAB>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.
J. Bacteriol. 183, 4823-4838, 2001
A;Title: Genome Sequence and Comparative Analysis of the Solvent-Producing Bacterium A;Reference number: A96900; MUID:21359325; PMID:21359325
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C;Superfamily: thioredoxin; thioredoxin homology
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A;Molecule type: DNA
A;Residues: 1-105 <KUR>
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C;Accession:
                                                                                           A; Description: PCR cloning
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I.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.
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    A;Cross-references:
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A;Cross-references: UNIPROT:P56430; UNIPARC:UPI0000136E82; GB:AE000594; GB:AE000511; NIE A;Experimental source: strain 26695
R;Alm, R.A.; Ling, L.S.L.; Moir, D.T.; King, B.L.; Brown, E.D.; Doig, P.C.; Smith, D.R., Ives, C.; Gibson, R.; Merberg, D.; Mills, S.D.; Jiang, Q.; Taylor, D.E.; Vovis, G.F.; Nature 397, 176-180, 1999
A;Title: Genomic sequence comparison of two unrelated isolates of the human gastric path A;Reference number: A71800; MUID:99120557; PMID:9923682
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C;Accession: H64622; C71890
R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, R. Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; McKen son, J.D.; Kelley, J.M.; Cotton, M.D.; Weldman, J.M.; Fujii, C.; Bowman, C.; Watthey, Nature 388, 539-547, 1997
A;Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, A;Title: The complete genome sequence of the gastric pathogen Helicobacter pylori.
A;Reference number: A64520; MUID:97394467; PMID:9252185
A;Accession: H64622
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                                                                                                                                                 C;Date: 11-Aug-1995 #sequence_revision 11-Aug-1995 #text_change 05-Oc C;Accession: A49888; S38886
C;Accession: A49888; S38886
R;Cohen, G: Argaman, A:, Schreiber, R.; Mislovati, M.; Aharonowitz, J. Bacteriol. 176, 973-984, 1994
A;Title: The thioredoxin system of Penicillium chrysogenum and its po
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C;Superfamily:
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                        A; Molecule type: DNA
A; Residues: 1-106 < COH>
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A; Residues: 1-106 < ARN>
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A; Residues: 1-106 < TOM>
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A;Variety: strains J99, 26695
                                                                      A; Status: preliminary
                                                                                                   A; Accession: A49888
                                                                                                                         A; Reference number: A49888;
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Best Local S
Matches 6
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Pred. No. 1.1;
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EMBL: X76120; NID: g426468;
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UNIPROT: P34723; UNIPARC: UPI0000136E91;

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A;Molecule type: protein
A;Residues: 1-107 <JOH>
A;Residues: 1-107 <JOH>
A;Cross-references: UNIPROT:P09857; UNIPARC:UPI0000136E76
A;Note: unidentified residues are Ile or Leu
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: heat-stable protein; redox-active disulfide
                                                                                                                                                                                                                                                 Biochemistry 26, 1209-1214, 1987
A;Title: The primary structure of thioredoxin from Chromatium
A;Reference number: A26622; MUID:87185419; PMID:3567166
A;Accession: A26622
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Gene: trxA
A;Introns: 8/2
C;Superfamily: thioredoxin; thioredox
F;10-92/Domain: thioredoxin homology
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C;Date: 31-Mar-1988 #sequence_revision 26-May-1994 #text_change 05-Oct-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, W.; Karp, P.; Romero, P.; Zhang, S.; Karp, P. 2317-2323, 2001
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A;Experimental source: strain C58 (Dupont)
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                                                                                                       10-93/Domain: thioredoxin homology <THR>
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Best Local Similarity
                                           Query Match
Local Similarity
nes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                       Chromatium vinosum
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100.0%; Score 47; DB 1; Length 107; llarity 100.0%; Pred. No. 1.1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                 redox-active #status predicted
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Pred. No. 1.1;
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M.; McClel
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thioredoxin C-1 [imported] - Brucella melitensis (strain 16M)
C;Species: Brucella melitensis
C;Date: 01-Feb-2002 #sequence_revision 01-Feb-2002 #text_change 05-Oct-2004
C;Accession: AH3504
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A;Residues: 1-107 <PYL>
A;Cross-references: UNIPROT:Q25345; UNIPARC:UPI000007DADA; GB:AE001274; NID:g3264850; PI
A:Experimental source: strain MHOM/IL/81/Friedlin
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A;Title: Leishmania major Friedlin chromosome 1 has an unusual distribution of protein-c A;Reference number: A81455; MUID:99178987; PMID:10077609
A;Accession: B81458
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C; Superfamily: thioredoxin; thioredoxin homology
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C;Accession: B61458; T02814
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C;Keywords: redox-active disulfide
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A; Residues: 1-107 < SAL>
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C;Species: Drosophila melanogaster
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change
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100.0%; Pred. No. 1.1;
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Pred. No. 1.1;
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C;Species: 1
C;Date: 18-1
C;Accession
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A;Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Venter, A;Title: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd. A;Reference number: A64000; MUID:95350630; PMID:7542800
A;Accession: E64047
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E64047
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A;Title: The genome sequence of the facultative intracellular pathogen Brucella melitens A;Reference number: AD3252; PMID:11756688
A;Title: Complete genome sequence of Pseudomonas aeruginosa PA01, an opportunistic pathon A;Reference number: A82950; MUID:20437337; PMID:10984043
A;Accession: G82991
A;Status: preliminary
A;Molecule type: DNA
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F;32-35/Disulfide bonds: redox-active #status predicted
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C; Superfamily:
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                                                                                                                       R;Stover, C.K.; Pham, X.Q.; Erwin, adman, S.; Yuan, Y.; Brody, L.L.; () LOTY, S.; Olson, M.V.
Nature 406, 959-964, 2000
                                                                                                                                                                                                                          thioredoxin pA5240 [imported] - Pseudomonas aeruginosa (strain PAO1) C;Species: Pseudomonas aeruginosa (c;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-C;Accession: G82991
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A; Residues: 1-107 <TIGR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hioredoxin - Haemophilus influenzae (strain Rd KW20)
;Species: Haemophilus influenzae
;Date: 18-Aug-1995 #sequence_revision 18-Aug-1995 #text_change 05-Oct-2004
;Accession: E64047
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Best Local
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K.; Lim,
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A;Residues: 1-108 <STO>
A;Cross-references: UNIPROT:Q9X2T1; UNIPARC:UPI0000136E95; GB:AE004936; GB:AE004091; NID A;Experimental source: strain PAO1
C;Genetics:
A;Gene: trxA; PA5240
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          thioredoxin VC0306 [imported] - Vibrio cholerae (strain N16961 serogroup O1) C;Species: Vibrio cholerae C;Date: 18-Aug-2000 #sequence revision 20-Aug-2000 #rave change of control of contr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, R.A.; Gwinn, M.L.; Dodson, R.J. chardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S.; Qin, H.; Dragoi, I.; Sellers, L. R.R.; Mekalanos, J.J.; Venter, J.C.; Fraser, C.M.
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A;Molecule type: DNA
A;Residues: 1-108 <KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nature 406, 477-483, 2000
A;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae.
A;Reference number: A82035; MVID:20406833; PMID:10952301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Gene: trxA
C;Superfamily: thioredoxin; thioredoxin homology
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C;Genetics:
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C;Date: 02-Nov-2001 #sequence_revision 02-Nov-2001 #text_change 05-Oct-2004
                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPROT:Q9KV51; UNIPARC:UPI00000C2C6E; GB:AE004119; GB:AE003852; A;Experimental source: serogroup O1; strain N16961; biotype El Tor
                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
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100.0%; Pred. No. 1.1;
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100.0%; Score 47; DI
100.0%; Pred. No. 1.:
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A; Molecule type: DNA
A; Roseidues: 'MLHQQNNQHARLIPVELY', 1-109 < WAL>
A; Residues: 'MLHQQNNQHARLIPVELY', 1-109 < WAL>
A; Residues: 'MLHQQNNQHARLIPVELY', 1-109 < WAL>
A; ROSE-ferences: UNIPROT: P00274; UNIPARC: UPI0000165538; GB: K02845; NID: g147610; PIDN:
A; Note: the sequence represents translation from an upstream ATG triplet that seems not
R; Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; Riley, M.; Cc
A.; Rose, D.J.; Mau, B.; Shao, Y.
Science 277, 1453-1462, 1997
Science 277, 1453-1462, 1997
A; Title: The complete genome sequence of Escherichia coli K-12.
A; Reference number: A64720; MUID: 97426617; PMID: 9278503
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A;Title: Characterization of the rho genes of Neisseria gonorrhoeae and Salmonella typhi A;Reference number: A49917; MUID:94075245; PMID:8253691
A;Accession: A49917;
               A; Molecule type: DNA
A; Residues: 'MLHQQRNQHARLIPVELY', 1-109 <BLAT>
A; Cross-references: UNIPARC: UPI0000165338; CB:AE000454; GB:U00096; NID:g2367278;
                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: Genetic and physical analysis of the thioredoxin (trxA) A;Reference number: A91519; MUID:85155506; PMID:6099324 A;Accession: A91519
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Nucleic Acids Res. 20, 1424, 1992
A;Title: Cloning and sequence of thioredoxin
A;Reference number: S35497; MUID:92220625; PM
A;Accession: S35497
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                                                                                                           A; Status: nucleic acid sequence not shown; translation not shown
                                                                                                                                          A;Accession: H65181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;Cross-references: UNIPARC:UPI000003112A; GB:Z21789; NID:g49361; PIDN:CAA79851.1; Note: submitted to the EMBL Data Library, February 1993
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;Experimental source: strain LT2
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B.J.; Kushner, S.R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       [validated] - Escherichia coli (strain K-12)
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6; Conserv
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substrain MG1655
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A;Note: the sequence represents translation from an upstream ATG triplet that seems R;Daniels, D.L.; Plunkett III, G.; Burland, V.; Blattner, F.R. Science 257, 771-778, 1992
A;Title: Analysis of the Escherichia coli genome: DNA sequence of the region from 84
A;Reference number: S30660; MUID:92358234; PMID:1379743
                                R,Lim, C.J.; Geraghty, D.; Fuchs, J.A. J. Bacteriol. 163, 311-316, 1985
                                                                                                                                                      A; Molecule type: DNA
A; Residues: 1-109 < M
                                                                                                                                                                                                                                                                A;Title: Autogenous regulation of the gene for transcription termination factor rho A;Reference number: I54863; MUID:86223816; PMID:2423505 A;Accession: I54863
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA A; Residues: 'MLHQQRN
                                                                                                              A; Cross-references: UNIPARC: UPI000003112A;
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(Cross-references: UNIPARC:UPI0000165538; EMBL:M87049; NID:g836656; PIDN:AAA67582.1; PI
(Cross-references: UNIPARC:UPI0000165538; EMBL:M87049; NID:g836656; PIDN:AAA67582.1; PI
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                                                                                                                                                  1-109 <MAS>
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                                                                                                              GB:M12779; NID:g148067; PIDN:AAA24694.1;
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A;Title: Cloning and nucleotide sequence of the trxA A;Reference number: A91802; MUID:85234377; PMID:3891; A;Accession: A91802 r cne crxA gene PMID:3891733 PID

A; Experimental source: strain K12 A;Cross-references: UNIPARC:UPI000003112A; GB:M10424; NID:g147608; PIDN:AAA24533.1; A;Residues: A; Molecule type: DNA 1-109 <LIM> PID

Biosci. Rep. 4, 917-923, 1984
A;Title: Nucleotide sequence of the thioredoxin gene from A;Reference number: I52550; MUID:85123150; PMID:6098320
A;Accession: I52550 R; Hoeoeg, Escherichia coli

A;Residues: A;Cross-references: UNIPARC:UPI000003112A; GB:M26133; NID:g148065; PIDN:AAA24693.1; A;Status: preliminary; translated from GB/EMBL/DDBJ Molecule type: DNA 1-109 <HOE>

A; Reference number: A91236; A; Title: Thioredoxin. R;Holmgren, Biochem. 6, 475-484, 1968
e: Thioredoxin. 6. The amino acid sequence of the protein rence number: A91236; MUID:69079993; PMID:4883076 from Escherichia coli

PID

A; Molecule type: protein
A; Residues: 2-16, 'LV', 19-71, 'IG', 74-109 <HOL>
A; Cross-references: UNIPARC: UPI0000171E22 A; Experimental source: strain B

A; Title: Structure requirements for disulfide bridge sulfitolysis of oxidized Escherich J. Biochem. 223, 473-479, 1994 engineered sequence

A;Reference number: \$45671; MUID:94333336; PMID:8055916
A;Contents: annotation; chemical activity of wild type and engin
R;Katti, S.K.; LeMaster, D.M.; Eklund, H.
submitted to the Brookhaven Protein Data Bank, March 1990
A;Reference number: A50900; PDB:2TRX
A;Contents: annotation; X-ray crystallography, 1.68 angstroms, x
R;Katti, S.K.; LeMaster, D.M.; Eklund, H.
J. Mol. Biol. 212, 167-184, 1990
A;Title: Crystal structure of thioredoxin from Escherichia coli residues 2-109

A; Reference number: A58630; MUID: 90204538; PMID: 2181145 at 1.68 Angstroms resolu

ç 2.8

A;Contents: annotation; X-ray crystallography, 1.68 angstroms R;Holmgren, A.; Soderberg, B.O.; Eklund, H.; Branden, C.I.
Proc. Natl. Acad. Sci. U.S.A. 72, 2305-2309, 1975
A;Title: Three-dimensional structure of Escherichia coli thioredoxin-S-2 A;Reference number: A93800; MUID:75176930; PMID:1094461
A;Rontents: annotation; X-ray crystallography, 2.8 angstroms R;Dyson, H.J.; Gippert, G.P.; Case, D.A.; Holmgren, A.; Wright, P.E. submitted to the Brookhaven Protein Data Bank, January 1990
A;Reference number: A50871; PDB:1TRX
A;Contents: annotation; conformation by (1)H-NMR, reduced form, residues R;Dyson, H.J.; Gippert, G.P.; Case, D.A.; Holmgren, A.; Wright, P.E. angstrom

Biochemistry 29, 4129-4136, 1990
A;Title: Three-dimensional solution structure of the reduced A;Reference number: A58631; MUID:90298180; PMID:2193685 form of Escherichia 2-109 coli

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C;Accession: Arvill Dougan, G.; James, K.D.; Thomson, N.K.; rivarion, T.; Connerton, P.; Cronin, A.; Davis, P.; Davies, R.M.; Down S.; Moule, S.; O'Gaora, P.
Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.; Title: Complete genome sequence of a multiple drug resistant A;Title: Complete yenome sequence of a multiple drug resistant
                                                                                                                                                   thioredoxin [imported] - Salmonella enterica subsp. enterica serovar Typhi C;Species: Salmonella enterica subsp. enterica serovar Typhi A;Note: this species has also been called Salmonella typhi C;Date: 09-Nov-2001 #sequence_revision 09-Nov-2001 #text_change 05-Oct-2004 C;Accession: AF0922
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C;Species: Emericella nidulans, Aspergillus nidulans
C;Date: 22-No-1993 #sequence_revision 01-Sep-1995 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Description: is reduced with NADPH by thioredo ol-disulfide exchange reactions C;Superfamily: thioredoxin; thioredoxin homology
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A;Reference number: A66948; PDB:1XOB
A; Reference number: AB0502; A; Accession: AF0922
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C;Superfamily: thioredoxin; thioredoxin homology
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A;Title: Purification, properties and primary structure of
A;Reference number: S27053; MUID:93092975; PMID:1459127
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R;Jeng, M.F.; Campbell, A.P.; Begley, T.; Holmgren, A.; Case, D.A.; Wright, P.E.;
submitted to the Brookhaven Protein Data Bank, November 1995
A;Reference number: A66947; PDB:1XOA
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Best Local Similarity
Matches 6; Conser
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Dyson
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A;Molecule type: DNA
A;Residues: 1-110 <STO>
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                                                                               A;Gene: NMB1366
C;Superfamily:
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    Conservative
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A,Authors: Grandi, G.; Sun, L.; Smith, H.O.; Fraser, C.M.; Moxon, E.R.; Rappuoli, R.; V. A;Title: Complete genome sequence of Neisseria meningitidis serogroup B strain MC58. A;Reference number: A81000; MUID:20175755; PMID:10710307
A;Accession: C81090
A;Status: preliminary
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-110 <TET-
A;Cross-references: UNIPROT:09JYY9; UNIPARC:UPI00000C46C4; GB:AE002485; GB:AE002098; NII
A;Experimental source: serogroup B, strain MC58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.E.; Eisen, J.; Heidelberg, B.; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; F. n, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, Proc. Natl. Acad. Sci. U.S.A. 98, 4135-4141, 2001
A;Title: Complete Genome Sequence of Caulobacter crescentus.
A;Reference number: A87249; MUID:21173698; PMID:11259647
A;Accession: A87688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Caulobacter crescentus
C;Date: 20-Apr-2001 #sequence_revision 20-Apr-2001 #text_change
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C; Superfamily: thioredoxin; thioredoxin homology
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0; Mismatches
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aft, D.H.; Kolon
C.; Fraser, C.M.
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C;Accession: E81850
                           A;Cross-references: UNIPARC:UPI0000171E27
R;Mittard, V.; Blackledge, M.J.; Stein, M.; Jacquot, J.P.; submitted to the Brookhaven Protein Data Bank, May 1996
                                                                                                                                                                                                     A;Cross-references: UNIPARC:UPI000016DD37; EMBL:X78822; NID:g840740; PIDN:CAA55399.1; R;Decottignies, P.; Schmitter, J.M.; Dutka, S.; Jacquot, J.P.; Miginiac-Maslow, M. Eur. J. Biochem. 198, 505-512, 1991

Bur. J. Biochem. 198, 505-512, 1991

A;Title: Characterization and primary structure of a second thioredoxin from the greer A;Reference number: S16090; MUID:91249849; PMID:2040309
                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPARC:UPI0000171E26
R;Stein, M.; Hodges, M.; Jeanette, E.; Lancelin, J.M.; Jacquot, J.P.
submitted to the EMBL Data Library, April 1994
A;Description: Chlamydomonas reinhardtii thioredoxins I : cDNA and amino acid deduced A;Reference number: S54844
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                                                                                                                        A; Molecule type: protein A; Residues: 2-112 < MIG>
                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: mRNA
A; Residues: 1-113 <STF>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: protein A; Residues: 2-15 <STW>
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A; Residues: 1-113 <S
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A;Accession: S57775
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;Date: 27-Oct-1995 #sequence revision 21-Jan-1997 #text_change 05-Oct-2004
;Date: 27-Oct-1995 #sequence revision 25-4870
;Accession: S57775; S57799; S54868; S16090; S54870
;Stein, M.; Jacquot, J.P.; Jeannette, E.; Decottignies, P.; Hodges, M.; Lancelin, J.M.
lant Mol. Biol. 28, 487-503, 1995
lant Mol. Biol. 28, 487-503, 1995
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Pred. No. 1.1;
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                                                                                                                                       R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Hc Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998

A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complet A;Reference number: A70500; MUID:98295987; PMID:9634230

A;Accession: B70851
                                                                                                                                                                                                                                                                                                                                                                              probable trxC protein - Mycobacterium tuberculosis (strain H37RV)
C;Species: Mycobacterium tuberculosis
C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 05-Oct-2004
C;Accession: B70851
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A;Residues: 1-116 <COL>
A;Cross-references: UNIPROT:P52229;
A;Experimental source: strain H37Rv
                                                                                      A; Molecule type: DNA
                                                                                                              A; Status: preliminary; nucleic acid sequence not shown; translation not
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RESULT 43
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A;Title: (1)H, (13)C, (15)N-NMR resonance assignments of oxidized thioredoxin R spectroscopy and computer-assisted backbone assignment.
A;Reference number: A58618; MUID:95262711; PMID:7744070
                                                                                                                                                                 A; Introns:
C; Superfam:
                                                                                                                                                                                                        A;Gene: CESP:B0228.5
                                                                                                                                                                                                                                                 A;Cross-references: UNIPROT:Q09433; UNIPARC:UPI0000136D67; EMBL:U23168; PIDN:AAC38808.1; A;Experimental source: strain Bristol N2
                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                     A; Reference number: Z18324
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submitted to the EMBL Data Library,
The sequence of C. e
                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Leimbach,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein 80228.5 - Caenorhabditis elegans C;Species: Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: redox-active disulfide F;2-113/Product: thioredoxin h #status experimental F;15-98/Domain: thioredoxin homology <THR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Contents: annotation; conformation by (1)H, (13)C, C; Genetics:
                                                                                                                                                                                                                                                                                            A;Residues: 1-115 <LEI>
                                                                                                                                                                                                                                                                                                                                                  A;Accession: T29044
                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            F;37-40/Disulfide bonds: redox-active #status experimental
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                                                                                                                                                               Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                 Matches
                                                                                                                      Query Match
                                                                                                     Local
                                                                                                                                                                                    69/3; 93/3
38
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                                                                             l Similarity
6; Conserv
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                                      WCGPCK 6
                                                                             100.0%;
llarity 100.0%;
Conservative 0
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elegans
                                                                                                   Score 47; DB:
Pred. No. 1.1;
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Pred. No. 1.1;
                                                                                 Mismatches
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cosmid B0228
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UNIPARC: UPI0000165202;

GB:AL021426;

GB:AL123456;

NID

complete

genome

Holroyd,

ß

shown

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thioredoxin (trxA) homolog - Lyme disease spirochete C.Species: Borrelia burgdorferi (Lyme disease spirochete) C.Date: 13-Feb-1998 #sequence_revision 13-Feb-1998 #text_change 05-Oct-2004 C.Accession: E70107 R.Fraser, C.M.; Casjens, S.; Huang, W.M.; Sutton, G.G.; Clayton, R.; Lathigra, R.; Son, D.; Peterson, J.; Kerlavage, A.R.; Quackenbush, J.; Salzberg, S.; Hanson, M.; Bowman, C.; Garland, S.; Fujii, C.; Cotton, M.D.; Horst, K.; Roberts, K.; Hatch, P. Nature 390, 580-586, 1997 Nature 390, 580-586, 1997 A.Authors: Smith, H.O.; Venter, J.C. A.; Title: Genomic sequence of a Lyme disease spirochaete, Borrelia burgdorferi. A.; Reference number: A70100; MUID:98065943; PMID:9403685
                                                                                                                                                                                                                                                                                     C;Accession: S34812
R;Brugidou, C.; Marty, I.; Chartier, Y.; Meyer, Y.
Mol. Gen. Genet. 238, 285-293, 1993
A;Title: The Nicotiana tabacum genome encodes two cytoplasmic A;Reference number: S34812; MUID:93241165; PMID:8479434
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A;Gene: trxC
A;Gene: trxC
C;Superfamily: thioredoxin; thioredox
F;15-98/Domain: thioredoxin homology
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                              S
                                                                                                                                                                                                                                                                                                                                                                        C;Species: Nicotiana tabacum (common tobacco)
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change
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                                                                                                                                                                                                                                  A; Molecule type: DNA
A; Residues: 1-118 < BRU>
                                                                                                                                                                                                                                                                                                                                                                                                                  thioredoxin h2 - common tobacco
                                                                                                                                                                                                                                                                                                                                                                                                                                    S34812
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A; Residues: 1-117 < KLE>
                                                                                                                                                                                                                 A;Cross-references: UNIPROT:Q07090; UNIPARC:UPI0000136D63;
                                                                                                                                                                                                                                                                      A; Accession: S3481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:O51088; UNIPARC:UPI000005732A; GB:AE001119; GB:AE000783; NIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Status: preliminary; nucleic acid sequence not shown; translation
                                                                                                                                         17-99/Domain: thioredoxin homology <THR>
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Best Local (
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38
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                                 WCGPCK 6
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WCGPCK 43
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thioredoxin; thioredoxin homology
                                                                                                                                                            thioredoxin; thioredoxin homology
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                                                                     Conservative
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                                                                                    100.0%;
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                                                                                    Score 47;
Pred. No.
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                                                                       Mismatches
                                                                                                       DB 1; Length 118;
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S19498
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A;Cross-references: SGD:S0000679
A;Map position: 3R
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A; Introns: 33/2
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A;Accession: T39387
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A; Residues: 1-127 < FEL>
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                                                                                                                                                            RESULT 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Accession: S19498
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                                                                                                                                             B91218
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Best Local S
Matches 6
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Best Local S
Matches 6
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thioredoxin ii - fission yeast (Schizosaccharomyces C;Species: Schizosaccharomyces pombe C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #Scacession: T39387 C;Accession: T39387 R;Lyne, M.; Rajandream, M.A.; Barrell, B.G.; Xiang, submitted to the EMBL Data Library, January 1999
                                             thioredoxin homolog YCR083w - yeast (Saccharomyces cerevisiae) C;Species: Saccharomyces cerevisiae C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_chang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:094504; UNIPARC:UPI000006BF82; EMBL:AL035085; PIDN:CAA22681.
A;Experimental source: strain 972h-; cosmid c12D12
                                                                                                                                                                                                                                                                                                                                                                                                                                C; Superfamily: thioredoxin; thioredoxin homology
                                             #sequence_revision 10-Sep-1999 #text_change 05-Oct-2004
                                                                                                                                                                                                                                                                                                                                                  100.0%;
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Pred. No. 1.2;
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R; Feldmann, H.; Mannhaupt, G.; Vetter, submitted to the Protein Sequence Datab Database, March 1992

A;Cross-references: UNIPROT:P25372; UNIPARC: UPI00001376B7; EMBL: X59720; NID: g1907116; P3

C;Keywords: redox-active disulfide F;34-115/Domain: thioredoxin homology ;55-58/Disulfide bonds: redox-active #status predicted Superfamily: thioredoxin; thioredoxin homology 100.0%; 0 Score 47; DB: Pred. No. 1.2; Mismatches ^THR> DB 1; 0; Length 127; Indels **;** Gaps

0

thioredoxin 1 [imported] - Escherichia coli (strain 0157:H7, substrain RIMD C;Species: Escherichia coli
C;Species: Bscherichia coli
C;Date: 18-7ul-2001 #sequence_revision 18-Jul-2001 #text_change 05-Oct-2004
C;Accession: B91218

0509952)

DNA Res. 8, 11-22, 2001
A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli 0157:H7 and A;Reference number: A99629; MUID:21156231; PMID:11258796
A;Accession: B91218
A;Status: preliminary R;Hayashi, T.; Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; Yokoyama, K.; gasawara, N.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; Shinagawa, H. geno ი.

RESULT

47

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R;Theologis, A; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso, Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K.; ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 816-820, 2000

A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, B.; Kim, C. C.A.; Li, Y.; Liu, S.X.; Liu, S.X.; Liu, S.A.; Luros, J.S.; Maiti, R.; Marziali, Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallon, ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Reference and analysis of chromosome 1 of the plant Arabidopsis.
A;Reference number: A86141; MUID:21016719; PMID:11130712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probable thioredoxin F23H11.5 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 05-Oct-2004
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C;Superfamily: thioredoxin; thioredoxin homology
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                                                                                                                     A; Molecule type: DNA
A; Residues: 1-129 <STO>
                                                                                                                                                                                              A; Reference number: A86141; A; Accession: B96621
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C;Superfamily: thioredoxin; thioredoxin homology
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A;Residues: 1-127 <STO>
A;Cross-references: UNIPARC:UPI0000165538; GB:AE005174;
A;Experimental source: strain O157:H7, substrain EDL933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, D.J.; iller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, E.; Potamousis, K.; A, Nature 409, 529-533, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Escherichia coli
C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 05-Oct-2004
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                                                  A; Gene: F23H11.5
                                                                        C; Genetics:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Accession: C86064
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                                                                                               A;Cross-references: UNIPROT:Q9XIF4; UNIPARC:UPI000000BF0A; GB:AE005173; NID:g5080807;
                                                                                                                                                                     A; Status: preliminary
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     Map position:
Superfamily:
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     thioredoxin
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Pred. No. 1
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Pred. No. 1.2;
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thioredoxin c-1 [imported] - Agrobacterium tumefaciens (strain C58, Cereon) C;Species: Agrobacterium tumefaciens C;Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 05-Oct-2004 C;Accession: F97361
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RESULT 53
D71707
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T08084
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                                                                                                                                                                                                                                                                                                               A;Cross-references: UNIPARC:UPI0000164C06; GB:AJ235270; A;Experimental source: strain Madrid E
                                                                                                                                                                                                                                                                                                                                                                                                                           C;Accession: D71707

R;Andersson, S.G.E.; Zomorodipour, A.; Andersson, J.O.; Sicheritz-Ponten, T.; Alsmark, Nature 396, 133-140, 1998

A;Title: The genome sequence of Rickettsia prowazekii and the origin of mitochondria. A;Reference number: A71630; MUID:99039499; PMID:9823893

A;Accession: D71707
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RESULT 54
                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: DNA
A; Residues: 1-130 < AND >
                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary; nucleic acid sequence not
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A; Residues: 1-129 < PAT>
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N;Alternate names: thioredoxin homolog
C;Species: Chlamydomonas reinhardtii
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Best Local S
Matches 6
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Matches
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                                                                                                                                                                                                                                                                                              Genetics:
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                                                                                                              WCGPCK 6
                                                                                                                                                      100.0%; Score 47; DB 2; Length 130; llarity 100.0%; Pred. No. 1.2; Conservative 0; Mismatches 0; Indels
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Pred. No. 1.2;
Mismatches
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C;Accession: B96721
R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; Wh Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khay C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Mait Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
R;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A;Reference number: A86141; MUID:21016719; PMID:11130712
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R;Lim, C.J.; Sa, J.H.; Fuchs, J.A.
Biochim. Biophys. Acta 1307, 13-16, 1996
A;Title: Identification of a third thioredoxin gene from Corynebacterium nephridii. A;Reference number: S70356; MUID:96254056; PMID:8652661
A;Accession: S70356
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A;Map position: circular chromosome
C;Superfamily: thioredoxin; thioredoxin homology
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A.; Liu, F.; Wollam, C.; Allinger, M.;
Science 294, 2323-2338, 2001
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A;Residues: 1-133 <KUR>
A;Cross-references: UNIPROT:Q8UJA6; UNIPARC:UPI00000D1691;
  A;Cross-references:
                     A; Molecule type:
A; Residues: 1-148
                                                                                                                                                                                                                                                                                                                                                                                         probable thioredoxin T17F3.9 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 05-Oct-2004
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Matches 6
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Matches 6
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Residues: 1-145 <LIM>
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UNIPROT: Q9CAS1;
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100.0%; Pred. No.
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100.0%; Pred. No. 1.3;
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UNIPARC:UPI00000C0F3; GB:AE005173; NID:g6358805;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.2;
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                                                                                                                                                                                                                              S.; Khaykin, B
.S.; Maiti, R.;
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                                                                                                                                                                                                                                                                                                                          , O.; Alonso,
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RESULT 58
T09495
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A;Title: Two functional thioredoxins containg redox-senesitive A;Reference number: Z16342; MUID:96198089; PMID:8626422
A;Accession: T08086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          dynein light chain, 16K, outer arm - Cl
N;Alternate names: thioredoxin homolog
C;Species: Chlamydomonas reinhardtii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Genetics:
A;Gene: T17F3.9
A;Map position: 1
C;Superfamily: thioredoxin
                                                                                                                                                                                                                                           A,Status: translated from GB/EMBL/DDB.
A;Molecule type: mRNA
A;Residues: 1-177 <DUE>
                                                                                                                                                                                                                                                                                                              R;Duek, P.D.; Mora Garcia, S.; Wolosiuk, submitted to the EMBL Data Library, Octol A;Reference number: Z16696
                                                                                                                                                                                                                                                                                                                                                                 thioredoxin m - rape chloroplast
C;Species: chloroplast Brassica napus
C;Date: 20-Sep-1999 #sequence_revision
C;Accession: T09495
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A;Experimental source: strain 1132D
C;Comment: Seems to interact directly with the a dynein heavy chain.
C;Superfamily: thioredoxin; thioredoxin homology
                            á
                                                                                                                                  C;Superfamily: thioredoxin; thioredoxin home C;Keywords: chloroplast; redox-active disulf;79-162/Domain: thioredoxin homology <TXN>
                                                                                                                                                                                                                                                                                            A; Reference number: A; Accession: T09495
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A; Residues: 1-156 < PAT>
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                                                                                   Similarity
 WCGPCK 105
                                                                                                                                                      thioredoxin; thioredoxin homology loroplast; redox-active disulfide
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Pred. No. 1.5
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Pred. No. 1.3;
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Pred. No.
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1 20-Sep-1999
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RESULT

103

WCGPCK 108

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C;Comment: This enzyme has amino-terminal heterogeneity. Thioredoxin m activates malate C;Comment: Arg-145 and Gly-164 are both essential for thioredoxin to interact with thior (;Superfamily: thioredoxin; thioredoxin homology C;Keywords: chloroplast; redox-active disulfide F;1-67/Domain: transit peptide (chloroplast) #status predicted <TNP> F;68-171/Product: thioredoxin mb #status experimental <PRD> F;69-171/Product: thioredoxin mc #status experimental <PRD> F;70-171/Product: thioredoxin mc #status experimental <PRD> F;70-171/Product: thioredoxin mc #status experimental <PRD> F;82-155/Domain: thioredoxin homology <THR> F;104-107/Disulfide bonds: redox-active #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross references: UNIPROT:D07591; UNIPARC:UPI0000136B67; GB:X51462; GB:S84848; NID:g21 R;Maeda, K.; Tsugita, A.; Dalzoppo, D.; Vilbois, F.; Schurmann, P. Eur. J. Biochem. 154, 197-203, 1986
A;Title: Further characterization and amino acid sequence of m-type thioredoxins from sp A;Reference number: A91158; MUID:86108311; PMID:3510868
A;Accession: JT0023
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A;Description: Genomic sequence for Arabidopsis thaliana
A;Reference number: Z14208
A;Accession: T00893
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: protein
A; Residues: 68-82, 'G', 84-89, 'Q', 91, 'SE', 94, 'S', 96-127, 'T', 129-164, 'D', 166, 'S', 168, 'YQ'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;Species: Spinacia oleracea (spinach)
;Date: 30-Sep-1988 #sequence revision 13-Mar-1998 #text_change 05-Oct-2004
;Accession: $20496; JT0023; B20273
;Accession: $20496; JT0023; B20273
;Wedel, N.; Clausmeyer, S.; Herrmann, R.G.; Gardet-Salvi, L.; Schuermann, P.
lant Mol. Biol. 18, 527-533, 1992
;Title: Nucleotide sequence of cDNAs encoding the entire precursor polypep;
;Reference number: $20496; MUID:92163017; PMID:1536927
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                                                                                                        Score 47; DB 1; Pred. No. 1.5;
                                                                                   Mismatches
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                                                                                                                                    Length 181;
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hypothetical protein F14I3.8 [imported] - Arabidopsis thaliana (;Speciaes: Arabidopsis thaliana (mouse-ear cress) C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change C;Accession: E96539 R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kan
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A, Molecule type: mRNA
A, Rolecule type: mRNA
A, Rolecule type: mRNA
A, Residues: 1-182 < LEP>
A, Cross-references: UNIPROT: P29450; UNIPARC: UPI0000136DE9; EMBL: X63537; NID: g20906; A, Cross-references: UNIPROT: P29450; UNIPARC: UPI0000136DE9; EMBL: X63537; NID: g20906; A, Cross-references: UNIPROT: thioredoxin; thioredoxin homology
C; Keywords: chloroplast; redox-active disulfide
F; 74-74/Domain: transit peptide (chloroplast) #status predicted <TNP>
F; 83-167/Domain: thioredoxin f #status predicted <MAT>
F, 83-167/Domain: thioredoxin homology <THR>
F; 106-109/Disulfide bonds: redox-active #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cnioredoxin f precursor - garden pea
C;Species: Pisum sativum (garden pea)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
C;Accession: S20929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Mora-Garcia, S.E.F.; Rodriguez-Suarez, R.J.; Wolosiuk, R.A. submitted to the EMBL Data Library, August 1997 A;Reference number: Z16164 A;Accession: T07837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Title: Isolation, characterization and nucleotide sequence A; Reference number: S20929; MUID:92256804; PMID:1581563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Lepiniec, L.; Hodges, M.; Gadal, P.; Cretin, Plant Mol. Biol. 18, 1023-1025, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C; Superfamily: thioredoxin; thioredoxin homology C; Keywords: chloroplast; redox-active disulfide
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C;Date: 14-May-1999 #sequence_revision 14-May-1999 #text_change 09-Jul-2004
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RESULT 63
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llarity 100.0%;
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Pred. No. 1.6;
Mismatches
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Pred. No. 1.6;
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A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White,

#text_change

05-Oct-2004

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Alonso,

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probable M-type thioredoxin [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_chang C;Accession: F85044
R;anonymous. The Finance R;anonymous.
R;anonymous, The European Union Arabidopsis Genome Sequencin Nature 402, 769-777, 1999
A;Title: Sequence and analysis of chromosome 4 of the plant A;Reference number: A85001; MUID:20083488; PMID:10617198
A;Accession: P85044
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-186 <STO>
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Nature 408, 816-820, 2010

A; Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R. Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.

A; Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, A; Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, A; Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.

A; Reference number: A86141; MUID:21016719; PMID:11130712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Tettelin, H.; Nelson, K.E.; Paulsen, I.T.; on, J.D.; Umayam, L.A.; White, O.; Salzberg, nson, T.; Hickey, E.K.; Holt, I.E.
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A; Authors: Loftus, B.J.; Yang, F.; Smith, H.O.; Venter, J.C.; Dougherty, A; Title: Complete Genome Sequence of a virulent isolate of Streptococcus
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A; Residues: 1-185 < KUR>
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Matches
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Best Local S
Matches 6
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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S.L.; Lewis, M.R.;
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Radune, D.; Holtzapple,
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       A;Description: activates fructose 1,6-bisphosphatase during photosynthesis
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A;Cross-references: UNIPROT:P09856; UNIPARC:UPI0000136DEA; EMBL:X14959; NID:g21343; A;Note: part of this sequence was confirmed by protein sequencing R;Tsugita, A.; Maeda, K.; Schuermann, P. Biochem. Biophys. Res. Commun. 115, 1-7, 1983
A;Title: Spinach chloroplast thioredoxins in evolutionary drift. A;Reference number: A20273; MUID:83308759; PMID:6351859
A;Molecule type: protein A;Recession: A20273
A;Rocession: A20273
A;Rocession: A20273
                                                                                                                                                                                                          R,Michalowski, C.B.; Bonnert, 4....
submitted to the RMBL Data Library, June 1998
submitted to the RMBL Data Library, June 1998
A:Description: A cDNA for
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A;Note: this sequence has been revised in reference S04661
A;Note: this sequence has been revised in reference S04661
C;Comment: Thioredoxin f activates fructose 1,6-bisphosphatase during photosynthesis
C;Superfamily: thioredoxin; thioredoxin homology
C;Keywords: chloroplast; redox-active disulfide
F;1-77/Domain: transit peptide (chloroplast) #status predicted <TNP>
F;78-190/Product: thioredoxin f #status predicted <MATP>
F;19-175/Domain: thioredoxin homology <THR>
F;19-175/Domain: thioredoxin homology <THR

F;19-175/Domain: thioredoxin homology <THR

F;19-175/Domain: thioredoxin homology <THR

F;19-175/Domain: thioredoxin homology 
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R;Kamo, M.; Tsugita, A.; Wiessner, C.; Wedel, N.; Bartling, D.

Bur. J. Blochem. 182, 315-322, 1989

A;Title: Primary structure of spinach-chloroplast thioredoxin
A;Reference number: S04661; MUID:89289715; PMID:2737203

A;Accession: S04661

A;Molecule type: mRNA
A;Residues: 1-190 <KRM>
A;Residues: 1-190 <KRM>
A;Residues: 1-190 <KRM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              thioredoxin f precursor - spinach
C;Species: Spinacia oleracea (spinach)
C;Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPROT:Q9SEU8; UNIPARC:UPI0000001768; GB:NC_001268; NID:g7270675; C;Genetics:
A;Gene: AT4903520
A;Map position: 4
C;Superfamily: thioredoxin; thioredoxin homology
                                                                                                                                                                                                                                                                                                                                                                             thioredoxin f precursor - common ice plant C;Species: Mesembryanthemum crystallinum (common ic C;Date: 23-Jul-1999 #sequence_revision 23-Jul-1999 C;Accession: T12261
                                       A; Molecule type: mRNA
A; Residues: 1-191 < MIC>
                                                                                                                                                                                                     A; Reference number: Z17478
A;Cross-references:
                                                                                                                      A;Status: preliminary; translated from GB/EMBL/DDB.
                                                                                                                                                                  A; Accession: T12261
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                F;114-117/Disulfide bonds: redox-active #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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Best Local
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UNIPROT: 081332; UNIPARC: UPI0000136DE8; EMBL: AF069314; NID: g3202021;
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Pred. No. 1.6;
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99 #text_change
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R;Hoskins, J.A.; Alborn Jr., W.; Arnold, J.; Blaszczak, L.; Burgett, S.; DeHoff, B. e, R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; McAhren, Y, P.; Sun, P.M.; Winkler, M.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Superfamily: thioredoxin; thioredoxin homology C;Keywords: redox-active disulfide F;92-176/Domain: thioredoxin homology <TXN>
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                                                                                                                                                                                                                                                                                                                   R;Lyne, M.H.; Rajandream, M.A.; Barrell, B.G.; Xiang, Z.; Hunt, I.C.; Aves, S.J. submitted to the EMBL Data Library, September 1999
A;Reference number: Z21936
A;Accession: T40552
                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Schizosaccharomyces pombe
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
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                                                                                                                                                                   A;Gene: SPAC577.08c
A;Map position: 2
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A;Experimental source: strain 972h-; cosmid c577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 69
T40552
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A; Residues: 1-191 < KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   J. Bacteriol. 183, 5709-5717, 2001
A;Authors: Yang, Y.; Young-Bellido, M.; Zhao, G.; Zook, C.; Baltz, R.H.; Jaskunas,
A;Title: Genome of the Bacterium Streptococcus pneumoniae Strain R6.
A;Reference number: A97872; MUID:21429245; PMID:11544234
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                                                                                                                                                                                                                                                                1;Residues: 1-290 <LYN>
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Best Local Similarity
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                                     WCGPCK 6
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Pred. No. 2.2
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Pred. No. 1.6;
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Search completed: February 23, Job time : 32.5 secs

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RESULT 70

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R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.B.; Chen, Y.; Woo, I erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClel: Karp, P.; Romero, P.; Zhang, S. Science 294, 2317-2323, 2001

A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan, W.; Perry, M.; Gordon-Kamm, ster, E.W.

Bter, E.W.

A;Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58.

A;Reference number: AB2577; MUID:21608550; PMID:11743193

A;Accession: AH3011
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A; Residues: 1-322 < KUR>
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                                                                                                                                                                                                                                                                                                                      A;Cross-references: UNIPROT:Q8U9M8; UNIPARC:UPI0000164857; GB:AE008689; PIDN:AAL44510.1; A;Experimental source: strain C58 (Dupont)
                                                                                                                                                                                                                              A; Map position: linear chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Status:
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Best Local
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   63
                                                         1 WCGPCK 6
                                                                                                                                             Similarity
WCGPCK 68
                                                                                                               Conservative
                                                                                                            100.0%; Score 47; DB:
100.0%; Pred. No. 2.3;
tive 0; Mismatches
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Catarrhini; Hominidae;

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Na Nicaud S., Jaffe D., Fisher S., Lutfalla G., Dossat C., Segurens B.,

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A Anthouard V., Jubin C., Castelli V., Katinka M., Vacherie B.,

A Anthouard V., Jubin C., Cattolico L., Poulain J., De Berardinis V.,

Cruaud C., Duprat S., Brottier P., Coutanceau J.P., Gouzy J.,

A Cruaud C., Lardier G., Chapple C., McKernan K.J., McEwan P., Bosak S.,

Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,

A Kellis M., Volff JN., Guigo R., Zody M.C., Mesirov J.,

A Lindblad-Toh K., Birren B., Musbaum C., Kahn D., Robinson-Rechavi M.,

A Lindblad-Toh K., Birren B., Musbaum C., Kahn D., Robinson-Rechavi M.,

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Cruaud C., Schachter V., Quetier F., Saurin W., Scarpelli C.,

Wincker P., Lander E.S., Weissenbach J., Roest Crollius H.,

"Genome duplication in the teleost fish Tetraodon nigroviridis reveals

the early vertebrate proto-karyotype.",

Mature 431:946-957(2004).
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Bukaryota; Metazoa; Chordata; Cranitata; Vertebrata; Euteleostomi;

Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;

Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;

Tetradontoidea; Tetraodontidae; Tetraodon.
                                                                                                       preliminary data.
EMBL; CAABO1008161; CAF91215.
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-i- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry whi
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fugu rubrip
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    Gaps
    0
RESULT 3
QST936 HUMAN P
AC QST936;
DT Q1-FEB-2005 (T
DE Thioredoxin.
GN Name-TXN, ORFN
OS Homo sapiens (
QC Eukaryota; Met
QC Eukaryota; Met
QC Mammalia; Euth
QC Mammalia; Euth
QC MCBI TaxID=960
RN (1) G000005489
DR GO; GO:0006489
DR GO; GO:0006418
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OROSTAL
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Best Local S
Matches 6
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EMBL; AF065241; AAC17430.1; -; mRNA.
HSSP; P10599; IRET.

SNR; O60744; 1-84.
GO: GO:0006618; P:electron transporter act GO: GO:0006618; P:electron transport; IEA InterPro; IPR006662; Thioredox dom2.
InterPro; IPR012335; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
Williams S.;
Submitted (MAY-2005) to the E
EMEL; AL158158; CAI14067.1; -
SMR; Q57936; 1-85.
GO; GO:0005489; F:electron to
GO; GO:0006118; P:electron to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-AUG-1998 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
Thioredoxin delta 3 (Fra
Mame=TXN delta 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pfam; PF00085; 'Injureuca..., -.. PRINTS; PR00421; THIOREDOXIN, PROSITE; PS00194; THIOREDOXIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             060744 HUMAN
060744;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-AUG-1998
01-AUG-1998
                                                                                                                                                                  NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Redox-active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2
HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WCGPCK 6
                                                                                                                                                                                                                                                                          ORFNames=RP11-427L11.1-002;
                                                                                                                                                                                                               Eutheria;
                                                                                                                                                                                                                               Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 1
84 AA; 9321 MW;
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                                                                                                                             SEQUENCE.
                                                                                                                                                                                                                                                                                                                 (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  center.
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                                                                                                                                                                                                                                                         (Human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Homin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rel. 07, Created)
rel. 07, Last sequence up
rel. 26, Last annotation
(Fragment).
                                                                                                                                                                                                           Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                   29,
    transport; IEA
                         transporter activity;
                                                            EMBL/GenBank/DDBJ
-; Genomic_DNA.
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Last annotation update)
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; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 47;
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n thioredoxin.";
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2; Length 84;

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Indels

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Q9HMDO
AC Q0
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Best Local S
Matches 6
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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EMBL; AY596297; AAV46961.1; -; Genomic DNA.

GO; GO:0005489; F:electron transporter acti
GO; GO:0005489; P:electron transport; IEA.

InterPro; IPR006662; Thiored. A

InterPro; IPR006663; Thioredoxin.

InterPro; IPR006663; Thioredox dom2.

Pfam; PF00085; Thioredoxin; 1.
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; ThioredoxIn-like.
Pfam; PF00085; ThioredoxIn; 1.
PRINTS; PR00421; THIOREDOXIN.
PROSITE; PS00194; THIOREDOXIN; 1.
Name=trxA2; OrderedLocusNames=VNG2600G; Halobacterium salinarium (Halobacterium halobium). Archaea; Euryarchaeota; Halobacteria; Halobacteriales;
                                                                                 Thioredoxin.
                                                                                                        01-MAR-2001 (TrEMBLrel.
01-MAR-2001 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                             Q9HMDO HALSA PRELIMINARY;
Q9HMDO;
01-MAR-2001 (TrEMBLrel. 1
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PubMed=15520287; DOI=10.1101/gr.2700304;
Baliga N.S., Bonneau R., Facciotti M.T., Pan M., Glusman G.,
Beltsch E.W., Shannon P., Chiu Y., Weng R.S., Gan R.R., Hung P.,
Date S.V., Marcotte E., Hood L., Ng W.V.;
"Genome sequence of Haloarcula marismortui: a halophilic archaeon from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Haloarcula marismortui (Halobacterium marismortui)
Archaea; Euryarchaeota; Halobacteria; Halobacteria
Halobacteriaceae; Haloarcula.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-FEB-2005
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01-FEB-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q5V0J2_HALMA PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TIGRFAMs; TIGR01068; thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
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                                                                                                                                                                                                                                                       HALSA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9
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88 AA; 10011 MW;
                                                                                                                                                                                                                                                                                                                                                                WCGPCK
                                                                                                                                                                                                                                                                                                                                                                                                                 WCGPCK 6
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                                                                                                           16,
26,
                                                                                                        Created)
Last sequence update)
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Pred. No. 2.1;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 47; DB 2; Length 88; Pred. No. 2.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CFE9B8FAC1F69857 CRC64;
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                                                                                                                                                                                                                          89
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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Best Local S
Matches 6
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15-JUL-1998
13-SEP-2005
Thioredoxin
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  EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         THIRO
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SQUENCE PRODUCTION OF THE PRODUCT OF
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RA Ng W.V., Kennedy S.P., Mahairas G.G., Berquist B., Pan M.,

Shukla H.D., Lasky S.R., Baliga N.S., Thorsson V., Sbrogna J.,

Swartzell S., Weir D., Hall J., Dahl T.A., Welti R., Goo Y.A.,

Leithauser B., Keller K., Cruz R., Danson M.J., Hough D.W.,

A Leithauser B., Keller K., Cruz R., Danson M.J., Hough D.W.,

Lienbarger T.A., Peck R.F., Pohlschroder M., Spudich J.L., Dung K.-H.,

A lasm M., Freitas T., Hou S., Daniels C.J., Dennis P.P., Omer A.D.,

Behardt H., Lowe T.M., Liang P., Riley M., Hood L., DasSarma S.;

"Genome sequence of Halobacterium species NRC-1.";

Proc. Natl. Acad. Sci. U.S.A. 97:12176-12181(2000).

EMBL; AE005133; AAG20641.1; -; Genomic_DNA.

PIR; E84409; E84409.
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GO; GO:0006118; P:electron transport; IEA.
GO; GO:0006110; P:transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR006663; Thioredox dom2.
Pfam; PF00085; Thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Haverkamp T., Schwenn J.D.; Submitted (AUG-1996) to the EMBL/GenBank/DDBJ databases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thiocapsa roseopersicina.
Bacteria; Proteobacteria;
                                                                                                                                                                                                                                      This Swiss-Prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE [GENOMIC DNA].
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                                                                                                                                                                                                                                                                                            similarity).
SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                                                                                                                                                                                                                   FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulfide and catalyzes dithiol-disulfide exchange reactions (By
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AAB36882.1;
2TRX.
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                                                                                                                                                                                                                               entry is copyright. It is produced through
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Pred. No. 2.2;
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Q4XKV3;
13-SEP-2005
13-SEP-2005
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PROSITE; P
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InterPro;
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EMBL; CAAJ01005024; CAH82459.1; -; Genomic_DNA.

InterPro; IPR006662; Thiored.

InterPro; IPR006663; Thioredox dom2.

InterPro; IPR012336; Thioredoxin-like.

InterPro; IPR012335; Thioredoxin fold.

Pfam; PF00085; Thioredoxin; 1.
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13-SEP-2005 (TrEMBLrel 31, Last sequence update)
13-SEP-2005 (TrEMBLrel 31, Last annotation updat
Thioredoxin, putative (Fragment).
ORFNames=PC300021.00.0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hall N., Karras M., Raine J.D., Carlton J.M., Kooij T.W.A., Berriman M., Florens L., Janssen C.S., Pain A., Christophides Bames K., Rutherford K., Harris B., Harris D., Churcher C., Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.
Hall N., Karras M., Raine J.D.,
Berriman M., Florens L., Jansse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Electron transport; Redox-active center; TDISULFID 33 36 Redox-active
                                                                                                                                                                                                                                                                                                                                                                     PRINTS; PR00421; THIOREDOXIN. PROSITE; PS00194; THIOREDOXIN;
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Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
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_РLАСН
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CAUTION: The sequence shown here is derived EWBL/GenBank/DDBJ whole genome shotgun (WGS)
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Pro; IPR005746; Thioredoxin-like.
Pro; IPR012336; Thioredoxin-fold.
Pr00085; Thioredoxin, 1.
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QSLB12; BACFN RELIMINARY;
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13-SEP-2005 (TrEMBLrel.
13-SEP-2005 (TrEMBLrel.
13-SEP-2005 (TrEMBLrel.
Thioredoxin, putative (F.
QRFNames=PB000593.00.0;
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Science 307:82-86(2005).
                                                                                                                                                       gene expression.";
Science 307:1463-1465(2005).
EMBL; CR626927; CAH08710.1; -;
                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

PubMed=15746427; DOI=10.1126/science.1107008;

Cerdeno-Tarraga A.-M., Patrick S., Crossman L.C., Blakely G.,

Abratt V., Lennard N., Poxton I., Duerden B., Harris B., Quai

Barron A., Clark L., Corton C., Doggett J., Holden M.T.G., La

Line A., Lord A., Norbertczak H., Ormond D., Price C.,

Rabbinowitech E., Woodward J., Barrell B.G., Pathill J.;

"Extensive DNA inversions in the B. fragilis genome control v
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PRINTS; PR00421
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EMBL; CAA101000448; CAH94443.1; -; Genc
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
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PRINTS; PR00421; THIOREDOXIN; UNKNOWN_1.
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Hall N., Karras M.,
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Eukaryota; Alveolata;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteroides fragilis (strain ATCC 25285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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EMBL/GenBank/DDBJ whole genome shotgun
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IPR006663; Thioredox dom2.
IPR005746; Thioredoxin.
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Pred. No.
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RESULT 11
QBAB91_BACTN ID QBAB91_F
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OS Bacteroides th
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PubMed=15466707; DOI=10.1073/pnas.0404172101;
Kuwahara T., Yamashita A., Hirakawa H., Nakayama H., Toh H.,
Kuhara S., Hattori M., Hayashi T., Ohnishi Y.;
"Genomic analysis of Bacteroides fragilis reveals extensive Inversions regulating cell surface adaptation.";
Proc. Natl. Acad. Sci. U.S.A. 101:14919-14924 (2004).
EMBL; AP006841; BAD49920.1; -; Genomic DNA.
GO; GO:0005489; F:electron transporter activity; IEA.
GO; GO:0006118; P:electron transport; IEA.
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Q64RG1;
25-OCT-2004 (TrEMBLrel. 2)
                            OrderedLocusNames=BT0219;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
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RESULT 12
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GG; GO:0006118; P:electron transport; IEA.
GG; GO:0006810; P:transport; IEA.
InterPro; IPR001452; SH3.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredoxin.
InterPro; IPR006563; Thioredoxin.
InterPro; IPR006663; Thioredoxin.
InterPro; IPR006663; Thioredoxin.
InterPro; IPR006663; Thioredoxin; 1.
PRINTS; PR00452; SH3DOMAIN.
PRINTS; PR00452; SH3DOMAIN.
PRINTS; PR00452; THIOREDOXIN.
ITIGRPAMB; TIGR01068; thioredoxin; 1.
PROSITE; PR00154; THIOREDOXIN; 1.
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10-OCT-2003 (Rel. 4
10-MAY-2005 (Rel. 4
Thioredoxin (Trx).
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28-FEB-2003
10-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=A5969Var.B;
MEDLINE=21956528; PubMed=11959450;
Skamrov A.V., Feoktistova E.S., Gol'dman M.A.,
"Mycoplasma gallisepticum rpoA gene cluster.";
FEMS Microbiol. Lett. 208:281-285(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacteroidetes; Bacteroides, NCBI TaxID=818;
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                                                                                            Microbiology 149:2307-2316(2003).

-!- FUNCTION: Participates in various redox reactions treversible oxidation of its active center dithiol tand catalyzes dithiol-disulfide exchange reactions similarity).
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Bacteria; Firmicutes; Mollicutes; Mycoplasmataceae; Mycoplasmataceaeae; Mycoplasmataceaeae; Mycoplasmataceaeae;
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Chiang H.C., Hooper L.V., Gordon J.I.;
"A genomic view of the human-Bacteroides thetaiotaomicron symbiosis.";
                                                                                                                                                                                                                                                                             "The complete genome sequence gallisepticum strain R(low).";
                                                                                                                                                                                                                                                                                                          Papazisi L., Gorton T.S., Kutish G., Markha
Nguyen D.K., Swartzell S., Madan A., Mahair
"The complete genome sequence of the avian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE 99 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=22830409; PubMed=12949158; DOI=10.1099/mic.0.26427-0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=R (low)
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                         SIMILARITY: Belongs to the thioredoxin family. SIMILARITY: Contains 1 thioredoxin domain.
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PubMed=12663928; DOI=10.1126/science.1080029;
PubMed=12663928; Dong S., Carmichael L.K.,
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Pred. No. 2.4;
; Mismatches
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Mahairas G., Geary S.J.
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RESULT 13
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MEDLINE=96320025; PubMed=87
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Skamrov A.V., Bibilashvili
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MCLEOTIDE SEQUENCE.
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STRAIN=A5969;
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Skamrov A.V., Bibilashvili
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Mol. Biol. (Mosk.) 30:585-5
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InterPro; IPR002746; ThioredoxIn.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
InterPro; IPR011194; Thioredoxin_like.
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                                      Skamrov A., Feoktistova E., (
Submitted (PEB-2003) to the I
EMBL; L35043; AAF36768.1; -;
HSSP; P80579; 1NW2.
                                                                                                                                                                                                                                                                       MEDLINE=96320025; PubMed=8754006;
Skamrov A.V., Bibilashvili R.Sh.;
Skamrov alvertain A5969
"A physical map of Mycoplasma gallisepticum strain A5969
determination of its positions on certain genes.";
Mol. Biol. (Mosk.) 30:585-594(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Skamrov A.V., Bibilashvili R.S.; "Mycoplasma gallisepticum strain S6 genome contains three regions hybridizing with 16 S rRNA and two regions hybridizing with 23 S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=A5969;
MEDLINE=92038016; PubMed=1718781; DOI=10.1016/0014-5793(91)81106-I;
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Bacteria; Firmicutes; Mollicutes;
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PRINTS; PR00421; THIOREDOXIN.
ProDom; PD003679; Thioredoxin like;
TIGRFAMs; TIGR01068; thioredoxin; 1
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sen the Swiss Institute of Bioinformatics and the EMBL
European Bioinformatics Institute. There are no restrict
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9; AAP56983.1; ALT_INIT; Genomic_DNA
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32 Redox-active (By sin
15 S -> T (in Ref. 1).
74 V -> I (in Ref. 1).
86 R -> I (in Ref. 1).
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26,
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Last seq
Last ann
                                                                                                         Goldman M., Beabealashvilli R.; EMBL/GenBank/DDBJ databases.
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annotation
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RESULT
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A Dodson R.J., DeBoy R.T., Gwinn M.L., Nelson W.C., Haft D.H.,
A Hickey E.K., Peterson J.D., Durkin A.S., Kolonay J.F., Yang F.,
A Holt I.E., Umayam L.A., Mason T.M., Brenner M., Shea T.P.,
A Holt I.E., Umayam L.A., Mason T.M., Brenner M., Shea T.P.,
A Radune D., Vamathevan J.J., Khouri H.M., White O., Gruber T.M.,
A Ketchum K.A., Venter J.C., Tettelin H., Bryant D.A., Fraser C.M.,
The complete genome sequence of Chlorobium tepidum TLS, a
photosynthetic, anaerobic, green-sulfur bacterium.",
Thotosynthetic, anaerobic, green-sulfur bacterium.",
Tho
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Best Local S
Matches 6
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28-FEB-2003
28-FEB-2003
13-SEP-2005
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InterPro;
InterPro;
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13-SEP-2005 (Rel. 48,
Thioredoxin 1 (Trx-1)
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PROSITE; F
                                                     InterPro;
InterPro;
                                                                                                                                                       EMBL; AE006470;
HSSP; P23400; 11
TIGR; CT0785; -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC STRAIN=TLS / ATCC 49652 / DSM 12025; MEDLINE=22103685; PubMed=12093901; DOI=10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=trx1; OrderedLocusNames=CT0785; Chlorobium tepidum.
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PRINTS; PR00421; THIOREDOXIN.
ProDom; PD003679; Thioredoxin_like; 1.
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InterPro; IPRO0
                  Pfam; PF00085; Th
PRINTS; PR00421;
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                                erPro; IPR006662; Thiored.
erPro; IPR006663; Thioredox dom2.
erPro; IPR005746; Thioredoxin.
erPro; IPR012336; Thioredoxin-like.
erPro; IPR012335; Thioredoxin_fold.
n; PP00085; Thioredoxin.
                                                                                                                                                                                                                                                          European a
                                                                                                                                                                                                                                                      Swiss-Prot entry is copyright. It is produced through a collaboratic een the Swiss Institute of Bioinformatics and the EMBL outstation European Bioinformatics Institute. There are no restrictions on it as long as its content is in no way modified and this statement is no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHLTE
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o; IPR006663; Thioredox dom2.
o; IPR005746; Thioredoxin.
o; IPR012336; Thioredoxin-like.
o; IPR012335; Thioredoxin fold.
o; IPR012335; Thioredoxin-like.
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TIGR01068;
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llarity 100.0%;
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                                                                                                                                                                                              AAM72022.1;
                    THIOREDOXIN
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41, Last sequence up
48, Last annotation
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thioredoxin; 1.
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C2DFB42A6DBF3C38
                                                                                                                                                                                            Genomic_DNA
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to a disulfide
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RESULT 16
THIO_CHLCV
ID THIO_CHLCV
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ID Q6KIE7_MYCMO
ID Q6KIE7,
AC Q6KIE7,
DT 05-JUL-2
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Matches 6
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Best Local 9
                                                                                                                                          P52227;
01-OCT-1996
01-OCT-1996
10-MAY-2005
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DISULFID
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE;
Complete
                                                          Chlamydophila caviae. Bacteria; Chlamydiae;
                                                                                                 Name=trxA; Synonyms=trx; OrderedLocusNames=CCA00080;
                                                                                                                       Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "The complete genome and proteome of Mycoplasma Genome Res. 14:1447-1461(2004).
EMBL; AE017737; AAT27629.1; -, Genomic_DNA.
HSSP; P10599; 1AUC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jaffe J.D., Stange-Thomann N., Smith C., Decaption.
Butler J., Calvo S., Elkins T., FitzGerald M.G., Haf
Kodira C.D., Major J., Wang S., Wilkinson J., Nicol
Birren B., Berg H.C., Church G.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=163K / ATCC 43663;
PubMed=15289470; DOI=10.1101/gr.2674004;
Jaffe J.D., Stange-Thomann N., Smith C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
STRAIN=163K / ATCC 4
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05-JUL-2004
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NUCLEOTIDE SEQUENCE
                                  NCBI_TaxID=83557;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRINTS; PR00421; THIOREDOXIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF00085; Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GO; GO:0005489; F:electron transporter activity; IEA. GO; GO:0006118; P:electron transport; IEA. InterPro; IPR006662; Thiored. InterPro; IPR006663; Thioredox dom2. InterPro; IPR012336; Thioredox dom2. InterPro; IPR012336; ThioredoxIn-like. InterPro; IPR012335; ThioredoxIn-fold.
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Q9PJK3;
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DISULFID
SEQUENCE
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                                                                                       Chlamydia muridarum.
Bacteria; Chlamydiae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; L39892; AAB41348.1; -; Genomic_DNA.
EMBL; AE016994; AAP04832.1; -; Genomic_DN
HSSP; P80579; 1QUW.
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"A 28 KDa major immunogen of Chlamydia psittaci shares identity wi
Mip proteins of Legionella spp. and Chlamydia trachomatis-cloning
characterization of the C. psittaci mip-like gene.";
Microbiology 142:945-953(1996).
                                                                                                                                                              Thioredoxin (TRX)
                                                                                                                                                                                    16-OCT-2001
10-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRINTS; PR00421; THIOREDOXIN.
TIGRFAM6; TIGR01068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           examining the role of niche-specific genes Chlamydiaceae.";
STRAIN=MOPn
                     NUCLEOTIDE SEQUENCE
                                                               NCBI_TaxID=83560;
                                                                                                                                        Name=trxA; OrderedLocusNames=TC0826
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MEDLINE=97090413; PubMed=8936321;
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IPRO06663; Thioredox dom2.
IPRO05746; Thioredoxin.
IPRO12336; Thioredoxin-like.
IPRO12335; Thioredoxin_fold.
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A Read T.D., Brunham R.C., Shen C., Gill S.R., Heidelberg J.F.,
A White O., Hickey B.K., Peterson J.D., Utterback T.R., Berry K.J.,
A Bass S., Linher K.D., Weidman J.F., Khouri H.M., Craven B., Bowman C.
A Dodson R.J., Gwinn M.L., Nelson W.C., DeBoy R.T., Kolonay J.F.,
A McClarty G., Salzberg S.L., Eisen W.C., DeBoy R.T., Kolonay J.F.,
A McClarty G., Salzberg S.L., Bisen W.C., DeBoy R.T., Kolonay J.F.,
A McClarty G., Salzberg S.L., Bisen W.C., Praser C.M.,
T "Genome sequences of Chlamydia trachomatis MoPn and Chlamydia
pneumoniae AR39.";
T "Uncleic Acids Res. 28:1397-1406(2000).

L Nucleic Acids Res. 28:1397-1406(2000).

-i- FUNCTION: Participates in various redox reactions through the
reversible oxidation of its active center dithiol to a disulfide
and catalyzes dithiol-disulfide exchange reactions.

-i- SIMILARITY: Contains 1 thioredoxin domain.
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Best Local :
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TIGREAMS;
PROSITE; F
                                                                                                                                                                                                                                                   STRAIN=D/UW-3/Cx;
MEDLINE=99000809; PubMed=9784136; DOI=10.1126/science.282.5389.754;
Stephens R.S., Kalman S., Lammel C.J., Fan J., Marathe R., Aravind
Mitchell W.P., Olinger L., Tatusov R.L., Zhao Q., Koonin E.V.,
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                                                                                                                                           Chlamydia trachomatis.";
Science 282:754-759(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=trxA; OrderedLocusNames=CT539;
Chlamydia trachomatis.
Bacteria; Chlamydiae; Chlamydiales; Chlamydiaceae; Chlamydia.
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE
                         FUNCTION: Participates in various redox reactions through reversible oxidation of its active center dithiol to a disand catalyzes dithiol-disulfide exchange reactions.
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s; TIGR01068; thioredoxin; 1.
; PS00194; THIOREDOXIN; 1.
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Matches 6
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[2]

NUCLEOTIDE SEQUENCE OF 1-21.

STRAIN=ATCC 33530 / G-37;

STRAINE=94075230; PubMed=8253680;

MEDLINE=94075230; PubMed=8253680;

MEDLINE=94075230; PubMed=8253680;

MCCOPIASMA Genitalium ge
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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P47370; Q49453;
01-FEB-1996 (Rel
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                                                                                                                                                                                                        STRAIN=ATCC 33530 / G-37;

MEDLINE=96026346; PubMed=7569993;

Fraser C.M., Gocayne J.D., White O., Adams M.D., Clayton R.A., Fleischmann R.D., Bult C.J., Kerlavage A.R., Sutton G.G., Kell Fritchman J.L., Weidman J.F., Small K.V., Sandusky M., Fuhrmar Nguyen D.T., Utterback T.R., Saudek D.M., Phillips C.A., Merri Tomb J.-F., Dougherty B.A., Bott K.F., Hu P.-C., Lucier T.S., Peterson S.N., Smith H.O., Hutchison C.A. III, Venter J.C.; "The minimal gene complement of Mycoplasma genitalium."; Science 270:397-403(1995).
          ++
                                                              J. Bacteriol. 175:7918-7930(1993).
-!- FUNCTION: Participates in various reversible exidation of its active
                                                                                                                                                                                                                                                                                                                                                                                        Name=trxA; Synonyms=trx; OrderedLocusNames=MG124; Mycoplasma genitalium.
Bacteria; Firmicutes; Mollicutes; Mycoplasmatacea
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01-FEB-1996 (Rel.
10-MAY-2005 (Rel.
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PHCI-2DPAGE; 084544;
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STRAIN=ATCC 33530 /
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European Bioinformatics Institute. There are no restrictions of as long as its content is in no way modified and this statement
        similarity).
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SIMILARITY: Contains 1
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31 Redox-active (By similarity).
11197 MW; BPAE022A5C980BE1 CRC64;
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             the thioredoxin family thioredoxin domain.
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
                                                                                                                                                                                                                                                  pneumoniae.";
Nucleic Acids Res. 24:4420-4449(1996).
-!- FUNCTION: Participates in various
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MEDLINE-97105885; PubMed-8948633; DOI=10.1093/nar/24.22.4420;
Himmelreich R., Hilbert H., Plagens H., Pirkl E., Li B.-C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "The thioredoxin reductase system Microbiology 143:1933-1940(1997).
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STRAIN=ATCC 29342 / M129;
MEDLINE=97346047; PubMed=9202470;
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Bacteria; Firmicutes; Mollicutes;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-11ke.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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EMBL; U01796; AAD12321.1; ALT_INIT; Genomic_DNA.
PIR; G64213; G64213.
HSSP; P07591; 1FB6.
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                                                                                                                                                                                                                                                                                                                           'Complete sequence analysis of the genome of the
                                                                                                                                                                                        FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulfide and catalyzes dithiol-disulfide exchange reactions (By
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is; TIGR01068; thioredoxin; 1.

pS00194; THIOREDOXIN; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     [LARGE SCALE GENOMIC DNA] .
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Matches
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HSSP;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR00746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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STRAIN=972
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Submitted (OCT-1997) to
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Cho Y.-W., Shin Y.H., Kim
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                                                                                                                                                                                                                                                                                                                                                                                                       PubMed=11859360; DOI=10.1038/nature724;
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Pred. No. 2.4;
0; Mismatches
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BJ databases.
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GeneDB Spombe; SPAC7D4.07c; -.

GO; GO:0005737; C:cytoplasm; TAS.

GO; GO:0005032; P:response to reactive oxygen specific properties of the control of
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PRINTS; PR00421; THIOREDOXIN.
TIGREAMS; TIGR01068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
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ACT_SITE 29 29 Nucleophi
ACT_SITE 32 32 Nucleophi
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TRX1_YEAST
P22217;
01-AUG-1991
01-OCT-1996
10-MAY-2005
Thioredoxin
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Nature 415:871-880(2002).
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EMBL;
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                                                                                                                                                                                                                          Similarity 6; Conser
                                                                                                                                                                                             WCGPCK 6
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 (Rel. 19, Created)
(Rel. 34, Last sequence update)
(Rel. 47, Last annotation updat
I (TR-I) (Thioredoxin 2).
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Pred. No. 2.4
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Deprotonates C-terminal active
(By similarity).
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R., Purnelle B.,
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Duesterhoeft A., Entian K.-D., Floeth M., Goffeau A., Hebling U., Heumann K., Heuss-Neitzel D., Hilbert H., Hilger F., Kleine K., Koetter P., Louis E.J., Messenguy F., Mewes H.-W., Miosga T., Koetter D., Mueller-Auer S., Nentwich U., Obermaier B., Piravandi Pohl T.M., Portetelle D., Purnelle B., Rechmann S., Rieger M., Rinke M., Rose M., Scharfe M., Scherens B., Scholler P., Schwager Schwarz S., Underwood A.P., Urrestarazu L.A., Vandenbol M., Verhasselt P., Vierendeels F., Voet M., Volckaert G., Voss H., Wambutt R., Wedler E., Wedler H., Zimmermann F.K., Zollner A., Wambutt R., Wedler E., Wedler H., Zimmermann F.K., Zollner A.,
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Saccharomycetales; Saccharomycetaceae; Saccharomyces
NCBI_TaxID=4932;
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                                                                                                                       PROTEIN SEQUENCE OF 25-33.

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Hall D.E., baldesten A., Holmgren A., Reichard P.;

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Hu Y., Vannberg F., Weger J., Kramer J., Moreira D., Kelley
Zuo D., Raphael J., Hogle C., Jepson D., Williamson J., Cama
Gonzaga L., Vasconcelos A.T., Simpson A., Kolodner R., Harlo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=97313267; PubMed=9169871;
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Anaorge W., Benes V., Brueckner M.,
Duesterhoeft A., Entian K.-D., Floet
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STRAIN=S288c
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MEDLINE=91225027; PubMed=2026619;
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MEDLINE=91107668; PubMed=1988444;
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PubMed=1561834;
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                           Schwenn J
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                                                PubMed=3060034;
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nucleotide sequence
re 387:87-90(1997).
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l. Chem. 266:1692-1696(1991).
  n J.D., Krone F. PAPS reductase:
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FUNCTION, AND DOTS AND TSA2 DEPENDENCE ON THIOREDOXIN.

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"LMA1 binds to vacuoles at Sec18p (NSF), transfers

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                                                                                                                                                                                                                                           Grant C.M.; "Role of th
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LEVEL OF PROTEIN EXPRESSION.
PubMed=14562106; DOI=10.1038/nature02046;
PubMed=14562108; Huh W.-K., Bower K., How
PubMed=14562108; Huh W.-K., Weissman J.S.;
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                                                                                                                                                                                                                                                                           PubMed=11169096;
                                                                                                                                                                                                                                                                                                                                                    "Roles of the glutathione- and thioredoxin-dependent in the Escherichia coli and Saccharomyces cerevisiae
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RESULT 23
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OF Putative
GN Name=trx
OS Chlamydo
OC Bacteria
OX NCBI_Tax
RN [1]
RP NUCLEOTI
RC STRAIN=S

Putative thioredoxin.

Chlamydophila abortus Bacteria; Chlamydiae;

Chlamydiae;

Chlamydiales;

Chlamydiaceae;

Chlamydophila

Name=trxA; OrderedLocusNames=CAB082

STRAIN=S26/3; NUCLEOTIDE NCBI_TaxID=83555;

SEQUENCE

[LARGE

SCALE

GENOMIC DNA]

Q5L731 CHLAB PRELIMINARY; Q5L731; Q1-FEB-2005 (TEMBLrel. 29, C 01-FEB-2005 (TEMBLrel. 29, I 01-FEB-2005 (TEMBLrel. 29, I

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                                                                                                                                                                                                                                                                                                                                                                                    protons. It is involved in many cellular processes, including deoxyribonucleotide synthesis, repair of oxidatively damaged proteins, protein folding, sulfur metabolism, and redox homeostasis. Thioredoxin-dependent enzymes include phosphoadenosine-phosphosulfate reductase MET16, alkyl-hydroperoxide reductase DOT5, thioredoxin peroxidases TSA1 and TSA2, alkyl hydroperoxide reductase AHP1, and peroxiredoxin HYR1. Thioredoxin is also involved in protection against reducing stress. As part of the LMA1 complex, it is involved in the facilitation of vesicle fusion such as homeotypic vacuole and ER-derived COPII vesicle fusion with the Golgi. This activity does not require the redox mechanism.

-i- SUBUNIT: Part of the haterodimeric LMA1 complex together with the proteinase inhibitor PBI2. LMA1 binds to the ATPase SEC18.
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"Yeast thioredoxin. Amino-acid sequence
disulfide of thioredoxin I and II.";
Eur. J. Biochem. 23:328-335(1971).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=22923954; PubMed=14562095; DOI=10.1038/nature02026; Hulf W.-K., Falvo J.V., Gerke L.C., Carroll A.S., Howson R.W Weissman J.S., O'Shea E.K.; "Global analysis of protein localization in budding yeast."
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MISCELLANBOUS: Yeast has two cytoplasmic thior
TRX2, and one mitochondrial, TRX3.
SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
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                                                                                                                                                                                European Bioinformatics Institute. There are no
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mitochondrial, TRX3
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THIOREDOXIN.
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Thomson N.R., Yeats C., Bell K., Holden M.T.G., Bentley S.D.,
Livingstone M., Cerdeno-Tarraga A.-M., Harris B., Doggett J.,
Ormond D., Mungall K., Clarke K., Feltwell T., Hance Z., Sanders
Quail M.A., Price C., Barrell B.G., Parkhill J., Longbottom D.;
"The Chlamydophila abortus genome sequence reveals an array of
variable proteins that contribute to interspecies variation.";
Genome Res. 15:629-640(2005).
EMBL; CR848038; CAH63539.1; -; Genomic_DNA.
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Butler J., Elkins
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Bacteria; Firmicutes; Mollicutes; Entomo
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Wipat A., Carter N., Brignell C.S., Guy J.B., Piper K., Sanders Wipat A., Carter N., Brignell C.S., Guy J.B., Piper K., Sanders Wipat A., Carter N., Brignell C.S., Guy J.B., Piper K., Sanders Wipat A., Carter N., Brignell C.S., Guy J.B., Piper K., Sanders Wipat A., Carter N., Emmerson P.T., Harwood C.R.;
"The dnaB-pheA (256 degrees-240 degrees) region of the Bacillus "The dnaB-pheA (256 degrees-240 degrees) region of the Bacillus subtilis chromosome containing genes responsible for stress responsible for stress
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P14949, 007960; Q45687;

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10-MAY-2005 (Rel 47, Last
MEDLINE=98044033; PubMed=9384377; DOI=10.1038/36786;
Kunst F., Ogasawara N., Moszer I., Albertini A.M., Alloni G.,
Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert
Borriss R., Boursier L., Brans A., Brann M., Brignell S.C., E
Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chen N.-Y., Zhang J.-J., Paulus H.;

"Chromosomal location of the Bacillus subtilis aspartokinase II gene and nucleotide sequence of the adjacent genes homologous to uvrC and trx of Escherichia coli.";
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Bacillus subtilis.
Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacil
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Pred. No. 2.4
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IEA.
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ine pleuropneumonia
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       Carter N.M.,
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RA Entlan K.-D., Exrington J., Fabret C., Ferrari E., Foulger D.,
RA Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N.,
RA Ghim S.-Y., Glaser P., Goffeau A., Golightly E.J., Grandi G.,
RA Guiseppi G., Guy B.J., Haga K., Haiech J., Harwood C.R., Henaut A.,
RA Hilbert H., Holsappel S., Hosono S., Hullo M.-F., Itaya M.,
Jones L.-M., Joris B., Karamata D., Kashara Y., Klaerr Blanchard M.,
RA Jones C., Kobayashi Y., Koetter P., Koningstein G., Krogh S.,
RA Klein C., Kobayashi Y., Koetter P., Koningstein G., Krogh S.,
RA Kumano M., Kurita K., Lapidus A., Lardinois S., Lauber J.,
RA Kumano M., Kurita K., Lapidus A., Lardinois S., Lauber J.,
RA Kumano M., Kurita K., Lapidus A., Lardinois S., Mauel C.,
RA Medigue C., Medina N., Mellado R.P., Mizuno M., Moesti D., Nakai S.,
RA Moback M., Noone D., O'Reilly M., Ogawa K., Ogiwara A., Oudega B.,
RA Noback M., Noone D., O'Reilly M., Ogawa K., Ogiwara A., Oudega B.,
RA Prescott A.M., Presecan E., Pujic P., Purnelle B., Rapoport G.,
RA Prescott A.M., Presecan E., Pujic P., Purnelle B., Rapoport G.,
RA Rose M., Sadaie Y., Sato T., Scanlan E., Schleich S., Schroeter R.,
RA Rose M., Sadaie Y., Sato T., Scanlan E., Schleich S., Schroeter R.,
RA Rose M., Sadaie Y., Sato T., Scanlan E., Takagi T., Takahashi H.,
RA Takemaru K., Takeuchi M., Tamakoshi A., Tanaka T., Terpstra P.,
RA Tognoni A., Viari A., Wambutt R., Wedler E., Wedler F.,
RA Weitzenegger T., Winters P., Wipat A., Yamamoto H., Yamane K.,
RA Yashimbara H. Danchia A., Wanitawa H.-F., Zumstein E.,
RA Yashimbara H. Danchia A., Jashida K., Yoshikawa H.-F., Zumstein E.,
Subtilist; BG10348; trxA.

R InterPro; IPR006662; Thioredox.dom2.

R InterPro; IPR006663; Thioredoxin.

R InterPro; IPR012346; Thioredoxin-like.

R InterPro; IPR012335; Thioredoxin-fold.

R InterPro; IPR012335; Thioredoxin-like.

R InterPro; IPR01234; Thioredoxin-like.

R InterPro; IPR01234; Thioredoxin-like.

R Pfam; PP00085; Thioredoxin; 1.

RRINTS; PR00421; THIOREDOXIN.

R PrODom; PD003679; Thioredoxin_like; 1.

R PTORFAMS; TIGR01068; thioredoxin; 1.

R PROSITE; PS00194; THIOREDOXIN; 1.

R Complete proteome; Direct protein sequencing; Electron transport;

W Redox-active center; Transport.
                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; J03294; AAA87311
EMBL; Z75208; CAA9957
EMBL; Z99118; CAB14811
EMBL; X79976; CAA5630
EMBL; X99275; CAA6766
PIR; B37192; B37192.
HSSP; P23400; 1DBY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=96345629; PubMed=8755892; Graumann P., Schroeder K., Schmid R., "Cold shock stress induced proteins i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        removed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yoshikawa H.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUNCTION: Participates in various redox reactions throu reversible oxidation of its active center dithiol to a and catalyzes dithiol-disulfide exchange reactions. SIMILARITY: Belongs to the thioredoxin family. SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          s Swiss-Prot entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EN European Bioinformatics Institute. There are no resti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       complete genome sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                long
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; CAA99577.1;
; CAB14810.1;
; CAA56300.1;
; CAA67667.1;
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V-1994) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              178:4611-4619 (1996).
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he EMBL/GenBank/DDBJ databases.
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Genomic_DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gram-positive bacterium Bacillus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and this statement
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                restrictions on
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to a disulfide
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RESULT 27
TRX2_YEAST
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Best Local (
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01-AUG-1991
01-OCT-1996
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Tettelin H., Agostoni-Carbone M.L., Albermann K., Albers M.,
Arroyo J., Backes U., Barreiros T., Bertani I., Bjourson A.J.,
Brueckner M., Bruschi C.V., Carignani G., Castagnoli L., Cerdan E.,
Clemente M.L., Coblenz A., Coglievina M., Coissac E., Defoor B.,
Del Bino S., Delius H., Delneri D., de Wergifosse P., Dujon B.,
Durand P., Entian K.-D., Eraso P., Becribano V., Fabiani L.,
Fartmann B., Fercli F., Feuermann M., Frontali L., Garcia-Gonzalez
Garcia-Saez M.I., Goffeau A., Guerreiro P., Hani J., Hansen M.,
Habling U., Hernandez K., Heumann K., Hilger F., Hofmann B.,
Indge K.J., James C.M., Klima R., Koetter P., Kramer B.,
Lauquin G., Leuther H., Louis E.J., Maillier E., Marconi A.,
Martegani E., Mazon M.J., Mazzoni C., McReynolds A.D.K.,
                                                                                                                                                                                                                                                                                                                                                                                             "Analysis
Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Saccharomyces cerevisiae (Baker's yeast).

Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                        Song
                                                                                                                                                                                                                phosphoserine phosphohydrolase, Yeast 12:273-280(1996).
                                                                                                                                                                                                                           VII reveals 12 ORFs, including CCT, ADE3 and TR-I genes, homologue the yeast PMT and EF1G genes, of the human and bacterial electron-transferring flavoproteins (beta-chain) and of the Escherichia cophosphoserine phosphohydrolase, and five new ORFs.";
                                                                                                                                                                                                                                                                                               MEDLINE=97060019; PubMed=8904340; DOI=10.1002/(STCI)1097-0061(19960315)12:3<273::AID-YEA898>3.3.CO;2-T; Guerreiro P., Barreiros T., Soares H., Cyrne L., Maia e Silva A., Rodrígues Pousada C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-OCT-1996 (Rel. 34, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation updat
Thioredoxin II (TR-II) (Thioredoxin 1).
                                                                                                                                                                    NUCLEOTIDE SEQUENCE (LARGE SCALE MEDLINE=97313265; PubMed=9169869;
                                                                                                                                                                                                 Yeast
[5]
                                                                                                                                                                                                                                                                    "Sequencing of a 17.6 kb segment VII reveals 12 ORFs, including CC
                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=S288c
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=91225027;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TRX2
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6; Conserv
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Chem. 266:9194-9202(1991)
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225027; PubMed=2026619;
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RA Paoluzi S., Plevani P., Portetelle D., Portillo P., Potier S.,
RA Purmelle B., Rieger M., Riles L., Rinaldi T., Robben J.,
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RA Souciet J.-L., Steensma H.Y., Talla B., Thierry A., Vandenbol M.,
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RA Volckaert G., Wambutt R., Watson M.D., Weber N., Wedler H.,
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STRAIN=S288C;
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Zuo D., Raphael J., Hogle C., Jepson D., Williamson J., Cama
Zuo D., Raphael J., Harlo
                                                                                                                                                                                                                                                                                                                                  "LMA1 binds to vacuoles at Secish (Vam3p) Coll 193:1125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FUNCTION, AND LMA1 COMPLEX.

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"A new_antioxidant with alkyl hydroperoxide defense propertie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FUNCTION, AND SULFATE ASSIMILATION AND PubMed=3060034; DOI=10.1007/BF00408300; Schwenn J.D., Krone F.A., Husmann K.;
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PubMed=14562106; DOI=10.1038/nature0;
Ghaemmaghami S., Huh W.-K., Bower K.,
Dephoure N., O'Shea E.K., Weissman J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Delaunay A., Isnard A.D., To "H202 sensing through oxidations of 19:5157-5166(2000).
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Carmel-Harel O., Storz G.;
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through the reversible oxidation of its active
disulfide, accompanied by the transfer of 2 ele
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Escherichia coli and Sac
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Biophys. Acta 1641:145-156(2003)
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39:533-541(2001).
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ard A.D., Toledano M.B.;
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DQ6FND5_CANGA PRELIMINARY;
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Best Local S
Matches 6
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PROSITE; F
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JO ASHGO
Q755CQO ASHGO PRELIMINARY;
Q75CQO;
                                                                                                                                                                                                         Candida glabrata (Yeast) (Torulopsis glabrata).

Bukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; mitosporic Saccharomycetales; Candida.
                                                                                                                                                                                                                                                                                                                          05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Candida glabrata strain CBS138 chromosome K complete sequence.
OrderedLocusNames=CAGLOK00803g;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dietrich F.S., Voegeli S., Brachat S., Erech A., Gate Mohr C., Poehlmann R., Luedi P., Choi S., Wing R.A., Gaffney T.D., Philippsen P.;
"The Ashbya gossypii genome as a tool for mapping the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ashbya gossypii (Yeast) (Eremothecium gossypii).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Eremothecium.
NCBI_TaxID=33169;
STRAIN=ATCC 2001 / CBS 138;
PubMed=15229592; DOI=10.1038/nature02579;
Dujon B., Sherman D., Fischer G., Durrens
                                                                                                                                                                      Eukaryota; Fungi; Ascomycota;
Saccharomycetales; mitosporic
NCBI_TaxID=5478;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; ThioredoxIn.
InterPro; IPR012336; ThioredoxIn-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
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STRAIN=ATCC 10895;
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05-JUL-2004
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                                                                                                      NUCLEOTIDE SEQUENCE
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GO:0006118; P:electron transport; IEA.
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llarity 100.0%;
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                                                                                                          [LARGE SCALE GENOMIC DNA]
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Pred. No. 2.4;
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   Casaregola S.,
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.A., Flavier A.,
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RESULT 30
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                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 6
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PROSITE; PS00194;
Complete proteome
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Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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GO; GO:0005489; F:electron transporter acti
GO; GO:00066118; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
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Nature 430:35-44(2004).
DEBHA
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103 AA;
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Pred. No. 2.4;
Mismatches
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RESULT 30

OGBURG DEBHA

OGBURG DEBHA PRELIMINARY; PRT; 103 AA.

AC QGBURG;

DT 25-OCT-2004 (TREMBLrel. 28, Last sequence update)

DT 25-OCT-2004 (TREMBLrel. 28, Last annotation update)

DT 25-OCT-2004 (TREMBLrel. 28, Last annotation update)

DE Similar to sp[P22217 Saccharomyces cerevisiae YLR043c TRX1 thioredoxin

DE I.

GN OrderedLocusNames=DEHA0G051269;

OS Debaryomyces hansenii (Yeast) (Torulaspora hansenii).

OS Debaryomyces hansenii (Yeast) (Torulaspora hansenii).

OS CUKALYOLE; Pungi; Ascomycota; Saccharomycetacea; Debaryomyces.

OX NCBI TaxlD=4959;

OX NCBI Tax

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Matches 6
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Jones T., Scherer S., Agabian N.;

"Annotation of the Genome of Candida albicans.";

Submitted (APR-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; AACQ011000032; EAL00485.1; -; Genomic DNA.

GO; GO:0005489; F:electron transporter activity; IEA.

GO; GO:000548; P:electron transporter activity; IEA.

R GO; GO:000548; P:electron transport; IEA.

R InterPro; IPR006662; Thioredox dom2.

R InterPro; IPR006663; Thioredoxin.

R InterPro; IPR012336; Thioredoxin.-11ke.

InterPro; IPR012335; Thioredoxin fold.
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Name=TRX1, ORFNames=Ca019.7611; Candida albicans SC5314.

Eukaryota; Fungi; Ascomycota; Saccharomycetales; mitosporic Sacc NCBI_TaxID=237561;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-1ike.
InterPro; IPR012335; Thioredoxin fold.
                                                                                                                                                                                                                                                                                                                                                                                          PubMed=15123810; DOI=10.1073/pnas.0401648101; Jones T., Federspiel N.A., Chibana H., Dungan J., Magee B.B., Newport G., Thorstenson Y.R., Agabian Davis R.W., Scherer S.; The diploid genome sequence of Candida albicans." Proc. Natl. Acad. Sci. U.S.A. 101:7329-7334(2004).
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PRINTS; PR00421; THIOREDOXIN.
TIGRFAMS; TIGR01068; thioredoxin; 1.
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                                                                                                                              Pfam; PF00085; Thioredoxin; PRINTS; PR00421; THIOREDOXIN
                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
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O:0005489; F:electron transporter_acti
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                           6; Conserv
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                                                                                                PS00194; THIOREDOXIN; 1.
 WCGPCK 6
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                           100.0%; ilarity 100.0%; Conservative 0
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ci. U.S.A. 101:7329-7334(2004).
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                           Score 47; DB Pred. No. 2.4 D; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Saccharomycotina; Saccharomycetes; Saccharomycetales; Candida.
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                                                                                     4B78100038A53620 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      activity; IEA.
                                        2.4;
                                                       DB 2;
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                                                         Length 103;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Kalman S.,
N., Magee
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                              Indels
                                                                                                                                                                                                                                                                                                                                    Adegbola O.,
                                                                                                                                                                                                                                                                                                                     K.-W.,
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                            <u>,</u>
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                            Gaps
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RESULT 33

QSFLW1 LACAC

ID QSFLW1 LACAC

QSFLW1;

AC QSFLW1;

AC QSFLW1;

DT 10-MAY-2005 (TREMBLrel. 30, L2

DT 10-MAY-2005 (TREMBLY 2)

DE Thioredoxin reductase

COLUMN action red
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Q7P4WB FUSNV PRELIMINARY;
ID Q7P4WB;
AC Q7P4WB;
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Best Local S
Matches 6
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GO; GO:0005489; F:electron transporter ac: GO; GO:0006118; P:electron transport; IEA.
GO; GO:0006810; P:transport; IEA.
InterPro; IPR006662; Thioredox.
InterPro; IPR006663; Thioredox.
InterPro; IPR006663; Thioredox.
Pfam; PF00085; Thioredox.

1. **Thioredox.** Thioredox.** Thiored
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAR-2004 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Karpatral V., Ivanova N., Anderson I., Reznik G., Bhatta-Gardner W.L., Mikhailova N., Larsen N., D'Souza M., Walu-Haselkorn R., Overbeek R., Kyrpides N.;
Submitted (JAN-2003) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry w
                                               PubMed=15671160; DOI=10.1073/pnas.0409188102; Altermann E., Russell W.M., Azcarate-Peril M.A., B. Buck B.L., McAuliffe O., Souther N., Dobson A., Du Lick S., Hamrick A., Cano R., Klaenhammer T.R.; "Complete genome sequence of the probiotic lactic Lactobacillus acidophilus NCFM."; Proc. Natl. Acad. Sci. U.S.A. 102:3906-3912(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lactobacillus acidophilus.
Bacteria; Firmicutes; Lactobacillales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=ATCC 49256;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fusobacterium nucleatum Bacteria; Fusobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=FNV0931;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE.
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                  Proc. Natl. Acad. Sci. U.S.A. 102:3906-3912(2005)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRINTS; PR00421; THIOREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   preliminary data.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 100 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WCGPCK 35
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TIGR01068;
103 AA; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; thioredoxin;
11369 MW; 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EAA23714.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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Fusobacterales; Fusobacteriaceae;
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Last annotation updat
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Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 47; DB 2
Pred. No. 2.4;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in; 1.
4437B8AAD056D7C0 CRC64;
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., Walunas T.,
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                                                                                                                           acid bacterium
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                                                                                                                                                                                               Callanan M.,
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Best Local S
Matches 6
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Best Local S
Matches 6
                                                                                                                                                      J. Bacteriol. 184.2005-2018 (2002).

2 EMBL; AE009951; AAL94306.1; -; Genomic_DNA.

2 HSSP; P00274; 1T7P.

3 GO; GO:0005489; F:electron transporter activity; IEA.

R GO; GO:0006118; P:electron transport; IEA.

R InterPro; IPR006662; Thioredox.dom2.

R InterPro; IPR006663; Thioredoxin.

R InterPro; IPR00663; Thioredoxin.

R InterPro; IPR012335; Thioredoxin-1lke.

R InterPro; IPR012335; Thioredoxin-fold.
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GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
PEam; PP00085; Thioredoxin, 1.
PRINTS; PR00421; THIOREDOXIN.
TIGRPAM6; TIGR01086; thioredoxin; 1.
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QBH23;
QBH23;
Q1-JUN-2002 (TrEMBLrel. 21,
Q1-JUN-2002 (TrEMBLrel. 21,
Q1-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE;
Complete
SEQUENCE
                                                                                                                                                                                                                                                                                          Kapatral V., Anderson I., Ivanova N., Reznik G., Los T., Lykidis i Bhattacharyya A., Bartman A., Gardner W., Grechkin G., Zhu L., Vasieva O., Chu L., Kogan Y., Chaga O., Golteman E., Bernal A., Larsen N., D'Souza M., Walunas T., Pusch G., Haselkorn R., Fonstein M., Kyrpides N.C., Overbeek R.; "Genome sequence and analysis of the oral bacterium Fusobacterium ruccleatum strain ATCC 25586.";
                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=ATCC 25586;
MEDLINE=21886394; PubMed=11889109;
DOI=10.1128/JB.184.7.2005-2018.2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OrderedLocusNames=FN0093;
Fusobacterium nucleatum (subsp. nucleatum).
Bacteria; Fusobacteria; Fusobacterales; Fusobacteriaceae;
                                                                                                                                      Pfam; PF00085; Thioredoxin; 1. PRINTS; PR00421; THIOREDOXIN.
                                                                                                                          TIGRFAMs; TIGR01068; thioredoxin; 1.
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FUSNN
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                                                Similarity 6; Conser
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103 AA;
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                      WCGPCK 6
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                                                Conservative
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                                                                                                  11342 MW;
                                                         100.0%;
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26,
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Last annotation updat
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                                             Score 47; DB 2
Pred. No. 2.4;
0; Mismatches
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Pred. No. 2.4
0; Mismatches
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                                                                                                27E003A221A4C780 CRC64;
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                                                                       DB 2;
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                                                                                                                                                                                                                                                                   RESULT
THIO_BC
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RESULT 35
Q74KU7; LACJO
ID Q74KU7;
AC Q74KU7;
AC Q74KU7;
AC Q74KU7;
AC Q74KU7;
DT 05-UUL-2004 (TrEMBLrel. 27,
DE TALOENSINE FAILURS)

CO Lactobacillus johnsonii.
CO STRAIN=NCC 533;
RN 011Est A.-C., Zwahlen M.-C.
RA Pittet A.-C., Zwahlen M.-C.
RA Pittet A.-C., Zwahlen M.-C.
RA Pittet B., Mercenier A., XI.
RT "The genome sequence of the
RT Lactobacillus johnsonii NCC
RL PCO: Natl. Acad. Sci. U.S..
DR EMBL; AE017201; AA500472.1;
DR HSSE; P1059; IAUC.
DR GO; GO:0005108; F:electron
DR GO; GO:0005108; F:electron
DR GO; GO:0005108; F:electron
DR GO; GO:0005108; Thioree
DR InterPro; IPR005663; Thioree
DR InterPro; IPR012335; Thioree
DR InterPro; IPR
     CCCCCRRRRRRROCOCOGGEDDDDAID
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Best Local S
Matches 6
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GO; GO:0005418; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
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097680;
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EMBL; AE017201; AAS08472.1; -; Genomic_DNA.
HSSP, P10599; IAUC.
                                                                         Terashima H., Gotoh S., Yagi K., Mizoguchi T.;
"CDNA sequence of bowine thioredoxin.";
DNA Seq. 10:331-33(1999)
-!- FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulfi
                                                                                                                                                                        NUCLEOTIDE SEQUENCE [MRNA].

MEDLINE=20189621; PubMed=10727087;

MEDLINE=20189621; PubMed=10727087;
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Bos taurus
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NCBI_TaxID=9913;
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Mammalia; Eutheria;
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SUBCELLULAR LOCATION: Cytoplasmic
                           and catalyzes dithiol-disulfide similarity).
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(Rel. 40, Last sequence update)
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Laurasiatheria; Cetartiodactyla; Ruminantia;
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RESULT 37
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                                                                                                  NUCLEOTIDE SEQUENCE [MRNA].

MEDLINE=21576430; PubMed=11719593; DOI=10.1093/molehr/7.12.1159;
Lopata A., Sibson M.C., Enders A.C., Bloomfield K.L., Gregory M.

Trapani G.D., Perkins A.V., Tonissen K.F., Clarke F.M.;

"Expression and localization of thioredoxin during early implant in the marmoset monkey.";
                                                                                                                                                                                                           Callithrix jacchus (Common marmoset).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Euarchontoglires, Primates, Platyrrhini;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin_fold.
Pfan; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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                       SIMILARITY: Belongs to the thioredoxin fam SIMILARITY: Contains 1 thioredoxin domain.
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                                             FUNCTION: Participates in various redox reactions through reversible oxidation of its active center dithiol to a dis and catalyzes dithiol-disulfide exchange reactions.
SUBCELLULAR LOCATION: Cytoplasmic (By similarity).
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RESULT 38 밁 Ś Query Match Best Local S Matches 6 DISULFID SEQUENCE This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its differentially expressed in normal chicken embryo fibroblasts.";
J. Biol. Chem. 263:9607-9611(1988). Name=TXN; Gallus gallus (Chicken). 01-AUG-1988 13-SEP-2005 THIO CH P08629; CHICK SITE SITE ACT InterPro; InterPro; EMBL; AF353204; AAK HSSP; P10599; 1ERT. SMR; Q9BDJ3; 1-104. NUCLEOTIDE SEQUENCE MEDLINE=88257080; Po Thioredoxin 01-AUG-1988 PRINTS; PR00421; THIOREDOXIN.
TIGRFAMs; TIGR01068; thioredoxin; use as -Jones S.W., Luk K.-C. Archosauria; Aves; Eukaryota; Metazoa; Electron PROSITE; Pfam; "Isolation of a chicken thioredoxin NCBI_TaxID=9031; InterPro; InterPro; [nterPro; similarity).
SUBCELLULAR LOCATION: Cytoplasmic (By simi SIMILARITY: Belongs to the thioredoxin fam SIMILARITY: Contains 1 thioredoxin domain. FUNCTION: Participates in various redox reactions reversible oxidation of its active center dithiol and catalyzes dithiol-disulfide exchange reactions CHICK PF00085; Thioredoxin; 30 long 1 WCGPCK n transport; T 0 E 31 E 34 E 25 Similarity Chem. WCGPCK 35 PS00194; THIOREDOXIN; IPR006662; Thiored.
IPR006663; Thioredox dom2.
IPR005746; Thioredoxin.
IPR012336; Thioredoxin-like.
IPR012335; Thioredoxin fold. 104 as its Conservative 33 32 8 STANDARD; 6 AAK30295.1; 08, Created)
08, Last sequence update)
48, Last annotation updat PubMed=2838473; Neognathae; content is in no way modified Chordata; Craniata; Vertebrata; Euteleostomi; Neognathae; Galliformes; Phasianidae; Phasiani 11626 31 25 25 [MRNA] 34  $\omega$ 32 Redox-active center; 100 ₩. <u>.</u> 0 By similarity.
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Nucleophile (By Score 47; DB Pred. No. 2.5 0; Mismatches Redox-active (By s: 3E0F5243A04B8E81 Nucleophile (By similarity).

Deprotonates C-terminal active (By similarity). similarity).
Contributes to redox potential Contributes to redox potential similarity) mRNA. and CDNA 104 update) (By similarity). Rous clone. B DΒ Transport. (By similarity)
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RESULT 39
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InterPro; IPR012335; Thioredoxin_fold.
InterPro; IPR012335; Thioredoxin, 1.
Pfam; PF00085; Thioredoxin; 1.
PROSITE; PR00421; THIOREDOXIN; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
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                                                                                                                           This Swiss-Prot entry is copyright. It is produced through a collaboratio between the Swiss Institute of Bioinformatics and the EMBL outstagion the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is no
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Laurasiatheria; Perissodactyla;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 47;
Pred. No. 2
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                                                                       mRNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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Equidae; Equus.
                                                                                                                                                                                a collaboration
                                                                                                                                                                                                                                                                                                disulfide
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InterPro; IPR012336; Thioredo
InterPro; IPR012335; Thioredo
Pfam; PF00085; Thioredoxin; 1
PRINTS; PR00421; THIOREDOXIN:
TIGRFAMS; TIGR01068; thioredo
                                                                                               THIO HUMAN STANDARD; PRT; 104 AA. P1059; Q96KI3; 01-JUL-1989 (Rel. 11, Created) 01-DEC-1992 (Rel. 24, Last sequence update) 13-SEP-2005 (Rel. 48, Last annotation update) Thioredoxin (ATL-derived factor) (ADF) (Surfaprotein) (SASP).
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INIT_MET
ACT_SITE
ACT_SITE
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MEDLINE=89008454; PubMed=3170595;

Wollman E.E., D'Auriol L., Rimsky L.,

Wollman E.E., D'Auriol Dessarps F.;

Wingfield P., Graber P., Dessarps F.;

"Cloning and expression of a cDNA for
J. Biol. Chem. 263:15506-15512(1988).
                                                                                                                                                                                                                                                                                                                                            Name=TXN; Synonyms=TRDX, TRX, TRX1;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SITE
                    Submitted (JUN-2000)
                              TISSUE=Lens;
Reddy P.G.,
                                                                                              "ATL-derived factor (ADF), an IL-2 rethioredoxin; possible involvement of
                                                                                                                                                                                                                                                      Gene 102:221-228(1991).
                                                                                                                                                                                                                                                                  MEDLINE=91340156; PubMed=187444; Tonissen K.F., Wells J.R.E.; "Isolation and characterization
                                                                                                                                                                                                                                                                                                                      Homo.
NCBI_TaxID=9606;
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NUCLEOTIDE SEQUENCE
                                                     NUCLEOTIDE
                                                                                      receptor induction.";
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6; Conserv
                                                                         8:757-764 (1989).
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; TIGR01068; thioredoxin; 1.
PS00194; THIOREDOXIN; 1.
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                Bhuyan D.K.,
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104
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Thioredoxin_fold.
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                    EMBL/GenBank/DDBJ databases.
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Nucleophile (By similarity).

Nucleophile (By similarity).

Nucleophile (By similarity).

Deprotonates C-terminal active site Cys

(By similarity).
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Contributes to redox potential
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receptor/Tac
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                                                                                                                                                                                      human
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                                                                                                       Matsui H., Hamuro J.,
i H., Yodoi J.;
tor/Tac inducer homologous
                                                                                                                                                                                                                                                                  thioredoxin-encoding genes.";
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Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buctow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Blatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Stapleton M.J., Usdin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

A Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

A Hitling M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

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Cherch A., Schein J.E., Jones S.J.M., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human

"Generation sequences.",

Proc. Matl Acad Scri U.S. A 90.16890-16901(2002).
oxidized and reduced states of human thioredoxin.";
Structure 2:503-522(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rieder M.J., Livingston R.J., Daniels M.R., Montoya M.A., Chung M.-W., Miyamoto K.E., Nguyen C.P., Nguyen D.A., Poel C.L., Robertson P.D., Schackwitz W.S., Sherwood J.K., Witrak L.A., Nickerson D.A.; "NIEHS-SNPs, environmental genome project, NIEHS ESI5478, Department of Genome Sciences, Seattle, WA (URL: http://egp.gs.washington.edu)."; Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
                                                                             STRUCTURE BY NMR.
MEDLINE=95006318; PubMed=7922028;
Qin J., Clore G.M., Gronenborn A.
                                                                                                                                                                                                                                                                                                                                Forman-Kay J.D., Clore G.M., Dricoll P.C., Wingfield P., Richards F.M., Gronenborn A.M.;

"A proton nuclear magnetic resonance assignment and secondary structure determination of recombinant human thioredoxin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted
                                                                                                                                                                  Forman-Kay J.D., Clore G.M., Wingfield P., Gr
"High-resolution three-dimensional structure
human thioredoxin in solution.";
Biochemistry 30:2685-2698(1991).
                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=90057393; PubMed=2684271; Forman-Kay J.D., Clore G.M., Dric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plasma membranes.";
Biochem. Biophys. Res.
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Xu J.Y., Xu L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (JUL-2000) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Cloning, purification and thioredoxin.";
                                                                                                                                                                                                                                                   MEDLINE=91159399; PubMed=2001356;
                                                                                                                                                                                                                                                                       STRUCTURE BY NMR.
                                                                                                                                                                                                                                                                                                               Biochemistry
                                                                                                                                                                                                                                                                                                                                                                                                                                       STRUCTURE BY NMR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Identification of a thioredoxin-related protein associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Martin H., Dean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROTEIN SEQUENCE OF 1-14.
MEDLINE=91151337; PubMed=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A., Lou M.F.;
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                                                                                                                                                                                                                                                                                                               28:7088-7097(1989).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PubMed=1998498;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Commun. 175:123-128(1991)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           U.S.A.
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                                                                                                                                                                                                           Gronenborn A.M.; re of reduced rec
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EMBL; X77584; CAA54687.1; -; mRNA.

EMBL; X54539; CAA38410.1; J; Genomic_DNA.

EMBL; X54540; CAA38410.1; JOINED; Genomic_DNA.

EMBL; X54541; CAA38410.1; JOINED; Genomic_DNA.

EMBL; J04026; AAA74596.1; -; mRNA.

EMBL; J04026; AAA74596.1; -; mRNA.

EMBL; AY004872; AAF87085.1; -; mRNA.

EMBL; AY004872; AAF87085.1; -; mRNA.

EMBL; AY004872; AAF87085.1; -; mRNA.

EMBL; AF548001; AAR33187.1; -; mRNA.

EMBL; BC003377; AAH03377.1; -; mRNA.
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MEDILINE=9639719; PubMed=8805557; DOI=10.1016/S0969-2126(96)00079-2;
Weichsel A., Gasdaska J.R., Powls G., Montfort W.R.;
"Crystal structures of reduced, oxidized, and mutated human thioredoxins: evidence for a regulatory homodimer.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=96347359; PubMed=8736558; DOI=10.1016/S0969-2126(96)00065-2; Qin J., Clore G.M., Kennedy W.P., Kuszewski J., Gronenborn A.M.; "The solution structure of human thioredoxin complexed with its targ from Ref-1 reveals peptide chain reversal."; Structure 4:613-620(1996).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        X-RAY CRYSTALLOGRAPHY (2.1 ANGSTROMS) OF MUTANT ASN-60.
MEDLINE-98039128; PubMed-9369469; DOI=10.1021/bi9710048;
Andersen J.F., Sanders D.A., Gasdaska J.R., Weichsel A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRUCTURE BY NMR. MEDLINE=96347359;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        receptor TAC (II2R/P55).
SUBUNIT: Homodimer.
INTERACTION:
Q92905:COPS; NDEXp=5; IntAct=EBI-594644, I
SUBCELLULAR LOCATION: Cytoplasmic.
SIMILARITY: Belongs to the thioredoxin fam:
SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reversible oxidation of its active center dithiol to a dismand catalyzes dithiol-disulfide exchange reactions. FUNCTION: ADF augments the expression of the interleukin-2 receptor TAC (ILZR/P55).
                                                                                                                                                                                                                                                                                                                                                                                                                                                     European Bioinformatics Institute.
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 1ERU;
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nies P., Miginiac-Maslow M., Mollman E.;
hioredoxin reactivity-structure/function z
Blophys Res Commun. 173:1375-1381(1990)
X - ray; @=1-104.
NMR; A=1-104.
NMR; A=1-104.
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NMR; A=1-104.
NMR; B=1-104.
NMR; @=1-104.
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RESULT 41
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SWISS-ZDPAGE; P10599; HUMAN.
SWISS-ZDPAGE; P10599; 8006; IEF.
PHCI-ZDPAGE; P10599; -.
Siena-ZDPAGE; P10599; -.
Ensembl; ENSG00000136810; Homo sa
HGNC; HGNC:12435; TXN.
H-INVDB; HIX0008275; -.
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01-APR-1993
13-SEP-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            An G., Wu R.;

"Thioredoxin gene expression is transcriptionally up-regulated by retinol in monkey conducting airway epithelial cells.";

Biochem. Biophys. Res. Commun. 183:170-175(1992).

-I- FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulfide and catalyzes dithiol-disulfide exchange reactions.

-I- SUBCELLULAR LOCATION: Cytoplasmic (By similarity).

-I- SIMILARITY: Belongs to the thioredoxin family.
INIT MET
ACT_SITE
ACT_SITE
                                                                                                                                                                                                                                                                                                                                                   EMBL; M84643; AAA36921.1;
PIR; JS0667; JS0667.
HSSP; P10599; 1ERT.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
                                                                                                                                                                                                                                                                                                                              HSSP; P10599; 1ERT
SMR; P29451; 1-104.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This Swiss-Prot entry is copyright, It is produced through a cobetween the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict use as long as its content is in no way modified and this statuse as long as its content is in no way modified and this statuse as long as its content is in no way modified and this statuse as long as its content is in no way modified and this status as long as its content is in no way modified and this status was supported by the content in the content is the content in the content in the content in the content is the content in the 
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MEDLINE=92181438; PubMed=1543487;
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                                                                         PRINTS; PR00421; THIOREDOXIN,
TIGRPAMS; TIGRO1068; THIOREDOXIN; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
Electron transport; Redox-active cen
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
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3TRX; NMR; @=1-104.
4TRX; NMR; @=1-104.
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(Rel. 25, Last sequence update)
(Rel. 48, Last annotation update)
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  Redox-active center; Transport.

0 By similarity.
31 Nucleophile (By similarity).
34 Nucleophile (By similarity).
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Pred. No. 2.5;
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Matches 6
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NUCLEOTIDE SEQUENCE [MRNA].

MEDLINE=89251607; PubMed=2785919;

Tagaya Y., Maeda Y., Mitsui A., Kndo N., Matsui H., Hamuro J.,

Tagaya Y., Maeda Y., Mitsui A., Kndo N., Matsui H., Yodoi J.;

Brown N., Arai K.-I., Yokota T., Wakasugi H., Yodoi J.;

"ATL-derived factor (ADF), an IL-2 receptor/Tac inducer homologous thioredoxin; possible involvement of dithiol-reduction in the IL-2 receptor induction.";

receptor induction.";

EMBO J. 8:757-764(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P10639; Q9DBRO;
01-JUL-1989 (Rel. 11, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
Thioredoxin (ATL-derived factor) (ADF).
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                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=129/Sv; TISSUE=Liver;
MEDLINE=95137382; PubMed=7835695; DOI=10.1016/0378-1119(94)00707-Y;
Matsui M., Taniguchi Y., Hirota K., Taketo M., Yodoi J.;
"Structure of the mouse thioredoxin-encoding gene and its processed pseudogene.";
Gene 152:165-171(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA].
STRAIN=C57BL/6J; TISSUE-Pancreas;
MEDLLNE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ERRATUM, AND SEQUENCE REVISION.

MEDILINE=94244626; PubMed=9187776;

Tagaya Y., Maeda Y., Mitsui A., Kndo N., Matsui H.,

Brown N., Arai K.-I., Yokota T., Wakasugi H., Yodoi

EMBO J. 13:2244-2244(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Name=Txn; Synonyms=Txn1; Mus musculus (Mouse).
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Redox-active (By similarity)
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Contributes to redox potential
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Gojobori T.
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RA Klausherg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausher R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Allechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Allechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Cassant T.L., Scheetz T.E.,
RA Stapleton M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schemutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J
                                                                                                                                                                                                                                                                                                                 EMBL; X77585; CAA54688.1; -; mRNA.

EMBL; D21859; BAA04881.1; -; Genomic DNA.

EMBL; AK007537; BAB25096.1; -; mRNA.

EMBL; AK007790; BAB25256.1; -; mRNA.

EMBL; BC010756; AAH10756.1; -; mRNA.

PIR; JC4068; S04107.

HSSP; P10599; 1ERT.

SMR; P10639; 1-102.

SWISS-2DPAGE; P10639; MOUSE.

Ensembl; ENSMUSG00000050612; Mus musculus.
                      MGI; MGI:98874; Txnl.
GO; GO:0005739; C:mitochondrion; IDA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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"Analysis of the mouse transcriptome based on functional annotation of control of the mouse transcriptome based on functional annotation of the mouse transcriptome based on functional control of the mouse transcriptome t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the European
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLECTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUBCELLULAR LOCATION: Cytoplasmic.
SIMILARITY: Belongs to the thioredoxin
SIMILARITY: Contains 1 thioredoxin doma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and catalyzes dithiol-disulfide exchange reactions. FUNCTION: ADF augments the expression of the interleukin-2 receptor TAC (ILZR/P55).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                full-length cDNAs.";
420:563-573(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              long as its content
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   THIOREDOXIN; 1.
                                                                                                                                         Thiored.
Thioredox dom2.
Thioredoxin-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              is in no way modified and this
                                                                                                                                                                                                                                                                                                                            Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           family.
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RESULT 43
THIO_OPHHA
ID THIO_O
AC Q98TX1
DT 28-FEB
DT 28-FEB
DT 28-FEB
DT 13-SEP
GR Name=T
OS QPhiop
OC Lepido
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RN [1]
CC -i- FU
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Best Local S
Matches
                                       PRINTS; PRO0421; THIOREDOXIN.
PROSITE; PS00194; THIOREDOXIN,
Electron transport; Redox-acti
INIT MET 0 0 B
ACT_SITE 31 31 NA
ACT_SITE 34 34 NA
SITE 25 25 D
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Q98TX1;
28-FEB-2003
28-FEB-2003
13-SEP-2005
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CONFLICT
SEQUENCE
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INIT_MET
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                                                                                                                                                                                                                                               InterPro;
InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Name=TXN;
Ophiophagus
                                                                                                                                                                                                                                                                                                                                                                                                                                                          similarity).
-!- SIMILARITY: Belongs to the thioredoxin family.
-!- SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Thioredoxin
                                                                                                                                                                                InterPro; IPR012335; Thiore Pfam; PF00085; Thioredoxin;
                                                                                                                                                                                                                                                                                                                                                                   use as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (NOV-2000) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE-Venom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=8665;
                                                                                                                                                                                                                                                                                                                                                                                                        between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lepidosauria; Squamata;
                                                                                                                                                                                                                          InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 jee ₩.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lapidae; Elapinae;
                                                                                                                                                                                               L; AF321769; AAK09384.1; -; mRNA.
P; P10599; 3TRX.
erPro; IPR006662; Thiored.
erPro; IPR006663; Thioredox dom2.
erPro; IPR012336; Thioredoxin-like.
erPro; IPR012335; Thioredoxin_fold.
                                                                                                                                                                                                                                                                                                                                                                                      European
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and catalyzes dithiol-disulfide exchange reactions (By
                                                                                                                                                                                                                                                                                                                                                                                    Swiss-Prot entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the Ev European Bioinformatics Institute. There are no restr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30
                                                                                                                                                                                                                                                                                                                                                                 long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Venom gland;
Liu H., Zhang Y.;
equence of Ophiophagus hannah venom gland thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6
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                                                                                                                                             PS00194; THIOREDOXIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  104
                                                                                                                                                                                                                                                                                                                                                                 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hannah (King cobra) (Naja hannah).
Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
a; Squamata; Scleroglossa; Serpentes; Colubroidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; ilarity 100.0%; Conservative 0;
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99
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25
    32
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                                                                                                                                                                                                                                                                                                                                                                   its content
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41, Last sequence up
48, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         [MRNA]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ophiophagus
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                                                                                                                          Redox-active center;
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                                                            By similarity.
Nucleophile (B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 47; DB 1
Pred. No. 2.5;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Redox-active (By similarity)
S -> C (in Ref. 4; BAB25256)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             similarity).
Contributes to redox potential
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (By similarity). Contributes to redox potential value
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Deprotonates C-terminal active
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Nucleophile
                    Nucleophile (By similarity).
Nucleophile (By similarity).
Deprotonates C-terminal active
(By similarity).
      Contributes
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                                                                                                                                                                                                                                                                                                                                                                     modified
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                                                                                                                        Transport
    redox potential
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Length 104;
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similarity)
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CRC64;
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                                             site Cys
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Best Local S
Matches 6
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           Transport.
INIT MET
ACT_SITE
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                                                                                                                                                                                                                                                                                                                                                                                Lee K.S., Tang W.K., Cheung P.Y., S
Submitted (MAY-2000) to Swiss-Prot.
                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUB=Liver;
Yu G.W., Xu J.Y., Xu L., Cher
"The cloning and expression
Submitted (MAY-2001) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P82460; Q95JF9;
16-OCT-2001 (Rel. 40, Created)
28-FEB-2003 (Rel. 41, Last seq
13-SEP-2005 (Rel. 48, Last ann
                                                                                                            InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR005746; Thioredoxin.like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam, PF00085; Thioredoxin; 1.
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Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                              PRINTS; PR00421; THICTIGREAMS; TIGRO1068; PROSITE; PS00194; TH
                                                                                                                                                                                  SMR; P82460; 1-104
                                                                                                                                                                                                        EMBL; AF382821; AAK60272.1; -;
                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Erythrocyte;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=TXN;
                                                                                                                                                                                                                                removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE
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                                                                   Direct protein
                                                                                                                                                                                                                                                                                                                                                                                                                 PROTEIN SEQUENCE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9823;
                                                                                                                                                                                                                                                                                                                       FUNCTION: Participates in various redox reactions the reversible exidation of its active center dithiol than deathlyzes dithiol-disulfide exchange reactions. SUBCELLULAR LOCATION: Cytoplasmic (By similarity). TISSUE SPECIFICITY: Erythrocytes.
                                                                                                                                                                                                                                                                                               SIMILARITY: Belongs to the thioredoxin family. SIMILARITY: Contains 1 thioredoxin domain.
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Laurasiatheria;
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annotation update)
                                                                   Electron transport; Redox-active
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Nucleophile (
Nucleophile (i
Deprotonates (
(By similarity
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Redox-active (By similarity).
, 852B96CBEF85OAFB CRC64;
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Pred. No.
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leophile (By similarity) leophile (By similarity) rotonates C-terminal act similarity).
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RESULT 45
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A COCCUCATION OF THE PROPERTY 
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Matches 6
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SEQUENCE
                                                                                                         Transport.
ACT_SITE
ACT_SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Johnson R.S., Mathews W.R., Biemann K., Hopper S.;
"Amino acid sequence of thioredoxin isolated from rabbi
determined by tradem mass spectrometry.";
J. Biol. Chem. 263:9589-9597(1988).
-i- FUNCTION: Participates in various redox reactions to reversible oxidation of its active center dithiol to and catalyzes dithiol-disulfide exchange reactions.
-i- SIMILARITY: Belongs to the thioredoxin family.
-i- SIMILARITY: Contains 1 thioredoxin domain.
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Name=:x,n;
Oryctolagus cuniculus (Rabbit).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metazoa; Enarchontoglires; Glires; Lagomorpha; Leporid
                                                                                                                                                                                                    TIGRFAMS;
                                                                                                                                                                                                                                                                                                                                                                                            PIR; A28086; A28086.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     use as
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NCBI_TaxID=9986;
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                                                                                                                                                              Direct protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Swiss-Prot entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the El European Bioinformatics Institute. There are no rest
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Best Local S
Matches
             GO; GO:0004791; F:thioredoxin-disulfide re
GO; GO:0006118; P:electron transport; TAS:
InterPro; IPR006662; Thioredox dom2.
InterPro; IPR005663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
PRINTS; PR00421; THIOREDOXIN.
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01-JUL-1989
01-JUL-1989
13-SEP-2005
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE [MRNA].
Xie Z.H., Liu C.Z., He Y.H.,
Submitted (OCT-2000) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OIHL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NIH - Mammalian Gene
Submitted (SEP-2003)
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodent
Muroidea; Muridae; Murinae; Rattus.
                                                                                                                                                                                       HSSP; P10599; 1ERT.
SMR; P11232; 1-103.
                                                                                                                                                                                                                     EMBL; X14878; CAA33019.1; -; mRNA.
EMBL; AP311055; AAG49923.1; -; mRN
EMBL; BCO58454; AAH58454.1; -; mRN
PIR; S04352; S04352.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE [MRNA].
                                                                                                                                                                                                                                                                                                                                                                                                   and catalyzes dithiol-disulfide exchange reactions similarity).
SUBCELIULAR LOCATION: Cytoplasmic (By similarity).
SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FUNCTION: Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disulf:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30
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11, Last
48, Last
   THIOREDOXIN; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            He Y.H., Wang A.M., Ma C. to the EMBL/GenBank/DDBJ
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                                                                                                                                                                     Rattus norvegicus.
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                                                                                                                                      activity;
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RESULT 47
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Best Local S
Matches
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SEQUENCE
                                                                                               InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
                                                                                                                                                        HSSP;
SMR; I
                                                                                                                                                                                                                       This Swiss-Prot entry is copyright. It is produced the Swiss Institute of Bioinformatics the European Bioinformatics Institute. There
                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chorda
Mammalia; Eutheria; Lauras
Pecora; Bovidae; Caprinae;
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                                                                           PROSITE;
                                                                                     PRINTS; PR00421; THIOREDOXIN
                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=95078463;
                                                                                                                                                                                                               36
                                                                                                                                                                                                                                                                     similarity).
SUBCELLULAR LOCATION: Cytoplasmic (By similarity).
SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                        P50413;
                                                                                                                                                                 Z25864; CAA81083.1;
P10599; 1ERT.
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                transport;
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                                                                                                                                                                                                                                                                                                                                                                       463; PubMed=7987015; Cleuter Y., Wollman
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aprinae; Ovis.
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25
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Nucleophile (By similarity).

Nucleophile (By similarity).

Deprotonates C-terminal active site C
(By similarity).

Contributes to redox potential value
                             By similarity.
Nucleophile (B)
Nucleophile (B)
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Pred. No. 2.5
0; Mismatches
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Contributes to redox potential
    Nucleophile (By similarity).
Nucleophile (By similarity).
Deprotonates C-terminal active
(By similarity).
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Redox-active (By
                                                                                                                                                                                                                                                                                                                                                                                                                                          Craniata; Vertebrata; Euteleostomi;
heria; Cetartiodactyla; Ruminantia;
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RESULT 48
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Query Match
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Matches 6
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Best Local Similarity
Matches 6; Conserv
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                                                                                                      TIGRFAMS;
PROSITE; P
                                                                                                                                                                                                                                                                                                This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use as long as its content is in no way modified and this statement is no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NÜCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

MEDLINB=21311952; PubMed=11418146; DDC1=10.1016/S0140-6736(00)04403-MEDLINB=21311952; PubMed=11418146; DDC1=10.1016/S0140-6736(00)04403-MEDLINB=21311952; PubMed=116, Submart H., Warayama A., Kurakami H., Hosoyama A., Canamori M., Matsumaru H., Maruyama A., Murakami H., Hosoyama A., Mizutani-Ui Y., Takahbashi N.K., Sawano T., Inoue R.-I., Kaito C., Mizutani-Ui Y., Takahbashi N.K., Sawano T., Inoue R.-I., Kaito C., Kanehisa M., Harakawa H., Kuhara S., Goto S., Yabuzaki J., Kanehisa M., Yamashita A., Oshima K., Furuya K., Yoshimo C., Shiba Hattori M., Ogasawara N., Hayashi H., Hiramatsu K.;

"Whole genome sequencing of meticillin-resistant Staphylococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            POAOK4; Q9ZEH4;
29-MAR-2004 (Rel. 43, Created)
29-MAR-2004 (Rel. 43, Last seq
10-MAY-2005 (Rel. 47, Last ann
                                                                                                                                                                                                                                                                                                                                                                                                                    Lancet 357:1225-1240(2001).

-I-FUNCTION: Component of the thioredoxin-thioredoxin reductase system. Participates in various redox reactions through the reversible oxidation of its active center dithiol to a disultant catalyzes dithiol-disulfide exchange reactions (By
                                                                                                                                               InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; ThioredoxIn.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam, PF00085; Thioredoxin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Staphylococcus aureus (strain Mu50 / ATCC 700699)
Bacteria; Firmicutes; Bacillales; Staphylococcus.
                                                                                                                                                                                                                                                        EMBL; BA000017; BAB57307.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thioredoxin (TRX)
                                                                                                                                     PRINTS; PR00421;
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                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY: Belongs to the thioredoxin family. SIMILARITY: Contains 1 thioredoxin domain.
                                                                                                                                                                                                                                                                                                                                                                                                              similarity).
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Similarity 6; Conserv
                                                        PR00421; THIOREDOXIN.
; TIGR01068; thioredoxin; 1.
PS00194; THIOREDOXIN; 1.
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104
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104
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Redox-active (By similarity).
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              Score 47;
Pred. No.
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Pred. No. 2
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Lancet 357:1225-1240(2001).
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Lencet 357:125-1250 (2001).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EW the European Bioinformatics Institute. There are no restruse as long as its content is in no way modified and this s
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RX PubMed=15213324; DOI=10.1073/pnas.0402521101;

RA Holden M.T.G., Feil E.J., Lindsay J.A., Peacock S.J., Day N.P.J.,

RA Holden M.T.G., Feil E.J., Lindsay J.A., Peacock S.J., Day N.P.J.,

RA Enright M.C., Foster T.J., Moore C.E., Hurst L., Atkin R., Barron A.,

RA Churcher C., Clark L., Corton C., Cromin A., Doggett J., Dowd L.,

RA Churcher C., Clark L., Corton C., Cromin A., Doggett J., Dowd L.,

RA Churcher C., Clark L., Corton C., Cromin A., Doggett J., Dowd L.,

RA Churcher C., Clark L., Corton C., Cromin A., Doggett J., Dowd L.,

RA James K.D., Lennard N., Line A., Mayes R., Moule S., Mungall K.,

RA James K.D., Lennard N., Line A., Mayes R., Moule S., Mungall K.,

RA Ormond D., Quail M.A., Rabbinowitsch E., Rutherford K.M., Sanders M.,

RA Sharp S., Simmonds M., Stevens K., Whitehead S., Barrell B.G.,

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RY "Complete genomes of two clinical Staphylococcus aureus strains:

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Staphylococcus aureus Bacteria; Firmicutes; NCBI_TaxID=282459;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
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Pfam; PF00085; Thioredoxin; 1.
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                                                               Name=trxA; OrderedLocusNames=SAS1079
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SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
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s; TIGR01068; thioredoxin; 1.
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PDA.OK6; QSZEH4;
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13-SEP-2005 (Rel. 48, I)
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PubMed=15213324; DOI=10.1073/pnas.04025211.01

Holden M.T.G., Feil E.J., Lindsay J.A., Peacock S.J., Day N.P.J.

Baright M.C., Foster T.J., Moore C.E., Hurst L., Atkin R., Barro.

Bason N., Bentley S.D., Chillingworth C., Chillingworth T.,

Churcher C., Clark L., Corton C., Cronin A., Doggett J., Dowd L.

Feltwell T., Hance Z., Harris B., Hauser H., Holroyd S., Jagels James K.D., Lennard N., Line A., Mayes R., Moule S., Mungall K.,

Ormond D., Quail M.A., Rabbinowitsch E., Rutherford K.M., Sander Sharp S., Simmonds M. Stevens Y. Whitehard T.
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                  Uziel O., Borovok I., Schreiber R., Cohen G., Aharonowit "Transcriptional analysis of the thioredoxin (trxA) and reductase (trxB) genes in Staphylococcus aureus."; Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                 Staphylococcus aureus.
Bacteria; Firmicutes;
NCBI_TaxID=1280;
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PROSITE; I
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SIMILARITY: Belongs to the thioredoxin fam.
SIMILARITY: Contains 1 thioredoxin domain.
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S; PR00421; THIOREDOXIN.
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6; Conserv
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IPRO06663; Thioredox dom2.
IPRO05746, Thioredoxin.
IPRO12336; Thioredoxin-like.
IPRO12335; Thioredoxin_fold.
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Participates
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
InterPro; IPR012335; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN;
ITCRFAMs; TIGRO1068; thioredoxin; 1.
PROSSITE; PS00194; THIOREDOXIN; 1.
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EMBL;
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                                                                   This Swiss-Prot entry is copyright. It is produced through a continuous the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC MEDLINE=22040717; PubMed=12044378; DOI=1 Baba T., Takeuchi F., Kuroda M., Yuzawa Nagai Y., Iwama N., Asano K., Naimi T.,
                                                                                                                                                                                                                                                                                                                                                               Staphylococcus aureus Bacteria; Firmicutes;
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renome and virulence determinants of high virulence community-
                                                                                                                                                                                                                                                                                                                                                                                              Name=trxA; OrderedLocusNames=MW1028;
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SIMILARITY: Belongs to the thioredoxin family.

SIMILARITY: Contains 1 thioredoxin domain.
                                                        36
                                                                                                                    similarity).
SIMILARITY: Belongs to the thioredoxin family.
SIMILARITY: Contains 1 thioredoxin domain.
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 BA000033;
P00274; 17
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32 Redox-active (By similarity)
11440 MW; FE61559DB27B3920 CRC64;
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Bacillales; Staphylococcus.
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8; DOI=10.1016/S0140-6736(02)08713-5;
Yuzawa H., Aoki K.-I., Oguchi A.,
imi T., Kuroda H., Cui L.,
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Complete DISULFID SEQUENCE

proteome, Electron transport, Red 29 32 Redox-active

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STEER REPRESENTATION OF THE PROPERTY OF THE PR
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin_fold.
InterPro; IPR012335; Thioredoxin_fold.
InterPro; IPR012335; Thioredoxin_fold.
InterPro; IPR012335; Thioredoxin; 1.
PRANTS; PR00421; THIOREDOXIN;
ITCRFAMs; TICRO1068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
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SEQUENCE
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DOI=10.1046/j.1365-2958.2003.03671.x;
DOI=10.1046/j.1365-2958.2003.03671.x;
Zhang Y.-Q., Ren S.-X., Li H.-L., Wang Y.-X., Fu G., Yang J.,
Zhang Y.-Q., Miao Y.-G., Wang W.-Y., Chen R.-S., Shen Y., Chen Z.,
Yuan Z.-H., Zhao G.-P., Qu D., Danchin A., Wen Y.-M.;
"Genome-based analysis of virulence genes in a non-biofilm-forming
Staphylococcus epidezmidis strain (ATCC 12228).";
Mol. Microbiol. 49:1577-1593(2003).
-I- FUNCTION: Component of the thioredoxin-thioredoxin reductase
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin_fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
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STRAIN=ATCC 12228;
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Bacteria; Firmicutes; Bacil
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HSSP; P00274; 17
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SIMILARITY: Belongs to the thioredoxin family.

SIMILARITY: Contains 1 thioredoxin domain.
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                                                                                                                                                                                                                                                                                                                           AA004435.1; -;
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43, Last sequence
48, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                    content is in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SCALE
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; FE61559DB27B3920 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 47; DB
Pred. No. 2.5;
                                                                                                                                                                                                                                                                                                                     Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            update)
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RESULT 56
Q6C399 YARLI
ID Q6C399 YARLI PRELIMINARY; PRT;
AC Q6C399;
DT 25-OCT-2004 (TrEMBLrel. 28, Created)
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RY PubMed=15229592; DOI=10.1038/natture02579;

RA Dujon B., Sherman D., Fischer G., Durrens P., Casaregola S.,

RA Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,

RA Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,

RA Goffard N., Frangeul L., Aigle M., Anthouard V., Babbur A., Barbe V.,

RA Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,

RA Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,

RA Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,

RA Hantraye F., Hemnequin C., Jauniaux N., Joyet P., Kachouri R.,

RA Hantraye F., Hemnequin C., Jauniaux N., Joyet P., Kachouri R.,

RA Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,

RA Nicaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,

RA Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,

RA Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,

RA Senion-Meyer M., Zivanovic Y., Bolottin-Fukhhara M., Thierry A.,

RA Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,

"Genome evolution in vesata".
                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 6
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nature 430:35-44(2004).

EMBL; CR382125; CAG99773.1; -; Genomic DNA.

GO; GO:0005489; F:electron transporter activity; IEA.

GO; GO:0006116; P:electron transport; IEA.

InterPro; IPR006662; Thiored.

InterPro; IPR006663; Thioredox dom2.

InterPro; IPR005746; Thioredoxin.

InterPro; IPR012336; Thioredoxin-like.

InterPro; IPR012336; Thioredoxin-fold.

PENNTS; PR00421; THIOREDOXIN.

PRINTS; PR00421; THIOREDOXIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kluyveromyces lactis (Yeast).
Bukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Kluyveromyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q6CN03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TIGRFAMs; TIGR01068; thioredox: PROSITE; PS00194; THIOREDOXIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Genome evolution in yeasts.";
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6; Conserv
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(TrEMBLrel. 28, Last sequence update)
(TrEMBLrel. 28, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            104 AA;
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                                                                                                                                                                                                                                                                                                                                                   Conservative
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                Score 47; DB 2; Length 104; Pred. No. 2.5; indels
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2.5;
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F., de Daruvar A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Weissenbach J.,
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Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
Barnay S., Blanchin S., Beckerich J.-M., Beyne E., Bleykasten C.,
Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
A Micaud J.-M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
Pellenz S., Potier S., Richard G.-F., Straub M.-L., Suleau A.,
Swennen D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
Zeniou-Meyer M., Zivanovic Y., Bolotin-Fukuhara M., Thierry A.,
Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
Wincker P., Souciet J.-L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 6
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GO; GO:0005489; F:electron transporter activi
GO; GO:000518; F:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
Submitted [2]
                                                                                                                                                                       Cryptococcus neoformans var. neoformans JEC21.
Eukaryota; Fungi; Basidiomycota; Hymenomycetes; Heterobasidiomycetes;
Tremellomycetidae; Tremellales; Tremellaceae; Filobasidiella.
NCBI_TaxID=214684;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-OCT-2004 (TrEMBLrel. 28, Last sequence update) 25-OCT-2004 (TrEMBLrel. 28, Last annotation updat Similar to sp|P22803 Saccharomyces cerevisiae YGF
                                                                                                                                                                                                                                                                                                               10-MAY-2005
Thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=CLIB 122 / E 150;
PubMed=15229592; DOI=10.1038/nature02579;
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Saccharomycetales; Dipodascace
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                                                     Van Aken S.,
                                                                                                                                                                                                                                                                                                                                                                                         Q5KK55;
10-MAY-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                         QSKKS5_CRYNE
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                                                                                                     STRAIN=JEC21;
                                                                                                                                                                                                                                                                                       ORFNames=CNC04200
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CRYNE
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TIGR01068; thioredoxin; 1.
                               (MAY-2004)
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                                                                             Amedeo P.,
                                                                                                                               SEQUENCE
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(TrEMBLrel. 30, 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
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                                                                             Roncaglia P., Vamathevan J., Utterback
                               the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                         Last sequence update)
Last annotation update)
2), putative.
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Pred. No. 2.5
0; Mismatches
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Saccharomycotina;
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cerevisiae YGR209c
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SEQUENCE

[LARGE

SCALE

GENOMIC DNA].

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QSSW38 CYNNE
QSSW38;
AC QSSW38;
DT 13-SEP-2
DT 13-SEP-
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Best Local S
Matches 6
          EMBL; AARYO1000013; EAL22161.1; -; Genomic DN GO; GO:0005489; F:electron transporter actIvi GO; GO:0006118; P:electron transport; IEA.

InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredoxin.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q55W38;
Q55W38;
Q55W38;
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
PRINTS; PR00421; THIOREDOXIN.
TIGRPAMB; TIGR01068; thioredoxin; 1.
                                                                                                                                                                                                                                                  STRAIN=B-3501À;
Fung E., Hyman R.W., Rowley D., Bruno D., Miranda M., Fuk
Wickes B.L., Fu J., Davis R.W.;
"Cryptococcus neoformans serotype D sequencing.";
Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry wh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Loftus B.J., Fung E., Roncaglia P., Rowley D., Amedeo P., Bruno D., Vamathevan J., Miranda M., Anderson I.J., Fraser J.A., Allen J.E., Bosdet I.E., Brent M.R., Chiu R., Doering T.L., Donlin M.J., D'Souza C.A., Fox D.S., Grinberg V., Fu J., Fukushima M., Haas B.J., Huang J.C., Janbon G., Jones S.J.M., Koo H.L., Krzywinski M.I., Kwon-Chung K.J., Lengeler K.B., Maiti R., Marra M.A., Marra R.E., Mathewson C.A., Mitchell T.G., Pertea M., Riggs F.R., Salzberg S.L., Schein J.E., Shvartsbeyn A., Shin H., Shumway M., Specht C.A., Syh B.B., Tenney A., Utterback T.R., Wickes B.L., Wortman J.R., Wye N.H., Kronstad J.W., Lodge J.K., Heitman J., Davis R.W.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cryptococcus neoformans var. neof
Eukaryota; Fungi; Basidiomycota;
Tremellomycetidae; Tremellales; T
NCBI_TaxID=283643;
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Science 307:1321-1324(2005).
                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE
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Pred. No.
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RESULT 60
Q9NPK9 PLAFA
ID Q9NPK9 PLAFA
AC Q9NPK9;
AC Q9NPK9;
DT 01-OCT-2000 (TrEMBLrel. 15,
DT 01-MAR-2004 (TrEMBLrel. 26,
DT 01-MAR-2004 (TrEMBLrel. 26,
DE Thioredoxin.
GN Name=trx;
GN Name=trx;
OS Plasmodium falciparum.
OC Eukaryota; Alveolata; Apicot
OX NCBI_TaxID=5833;
RN [1]
RP NUCLEOTIDE SEQUENCE.
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Best Local
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PROSITE; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GO; GO:0005489; F:electron transporter activity; IEA. GO; GO:0006118; P:electron transport; IEA. InterPro; IPR006662; Thiored. InterPro; IPR006663; Thioredox dom2. InterPro; IPR005746; Thioredoxin. InterPro; IPR005746; Thioredoxin-like. InterPro; IPR012336; Thioredoxin-fold. InterPro; IPR012335; Thioredoxin-fold. Pfam, PF00085; Thioredoxin, 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kanzok S.M., Schirmer R.H., Turbachova I., Iozef R., Becker K.; "The thioredoxin system of the malaria parasite Plasmodium falciparum Glutathione reduction revisited.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       J Biol Chem. 275:40180-40186 (2000).
EMBL; AF202664; AAF34541.1; -; mRNA.
HSSP; P80028; 1EPF.
SMR; Q9NIR2; 1-103.
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Q9NIR2;
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PROSITE; PS00194; THIOREDOXIN;

HYPOchetical protein.

SEQUENCE 104 AA; 11451 MW;
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MEDLINE=20568274; Po
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Eukaryota; Alveolata;
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                                                                         Apicomplexa; Haemosporida; Plasmodium
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Pred. No.
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2F507E17F276171F CRC64;
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Submitted (MAY-2000) to the EMBL/GenBank/DDBJ databases.
R EMBL; AJ277839; CAB90828.1; -; mRNA.
R GO; GO:0005489; F:electron transporter activity; IEA.
R GO; GO:0006118; P:electron transport; IEA.
R InterPro; IPR006662; Thiored.
R InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012336; Thioredoxin-fold.
Pfam; PF00085; Thioredoxin; 1.
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Best Local (
                                                                                     SMR; Q7RH10; 1-103.

GO; GO:0005489; F:electron transporter a GO; GO:0006118; P:electron transport; IE InterPro; IPR006662; Thiored.

InterPro; IPR006663; Thioredox dom2.

InterPro; IPR005746; Thioredoxin-like.

InterPro; IPR012336; Thioredoxin-like.

InterPro; IPR012335; Thioredoxin-fold.

Pfam; PF00085; Thioredoxin; 1.
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Q7RH10;
Q1-MAR-2004
Q1-MAR-2004
Q1-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=22255706; PubMed=12368865; DOI=10.1038/nature01099; Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M., Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L., Slerson J.D., Pop M., Kosack D.S., Shumway M.F., Bidwell S.L., Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V., Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V., Cho J.K., Quackenbush J., Sedegah M., Shoabib A., Cummings L.M., Florens L., Yates J.R. III, Raine J.D., Sinden R.E., Harris M.A., Cunningham D.A., Preiser P.R., Bergman L.W., Vaidya A.B., van Lin L.H., Janse C.J., Waters A.P., Smith H.O., White O.R., Salzberg S.L., Venter J.C., Fraser C.M., Hoffman S.L., Gardner M., Carlos J.L., Venter J.C., Fraser C.M., Hoffman S.L., Gardner M.
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Krnajski Z., Gilberger T.W.,
Submitted (MAY-2000) to the
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TIGREAMS; TIGRO1068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
SEQUENCE 104 AA; 11716 MW; 01299
                     PRINTS; PR00421; THIOREDOXIN.
TIGRFAMB; TIGR01068; thioredoxin;
PROSITE; PS00194; THIOREDOXIN; UNI
                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Genome sequence and comparative analysis parasite Plasmodium yoelii yoelii."; Nature 419:512-519(2002).
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Eukaryota; Alveolata; Api
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HSSP; P80028; 1EP7.
SMR; Q7RH10; 1-103.
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PDB; 1SYR; X-ray; A/B/C/D/E/F/G/H/I/J/K/L1-1-104.

GO; GO:0005489; F:electron transporter activity; IE
GO; GO:0066118; P:electron transport; IEA.

InterPro; IPR006662; Thiored.

InterPro; IPR006663; Thioredox dom2.

InterPro; IPR005746; Thioredoxin.

InterPro; IPR012336; Thioredoxin-like.

InterPro; IPR012336; Thioredoxin-fold.
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05-JUL-2004
Thioredoxin.
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Q7KQL8;
Q7KQL8;
05-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE.
MEDLINE=22255705; PubMed=12368864;
MEDCANTER M.J. Hall N., Fung E., Whi
Gardner M.J., Hall N., Pelson K.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Plasmodium falciparum (isolate Eukaryota; Alveolata; Apicomple NCBI TaxID=36329;
                      Name=Trx-2;
Schistosoma
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  Eukaryota; Metazoa;
Schistosomatoidea;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Nature
                                                                                                                                                                                                                                                                                                                                                                                                                               "Genome sequence of falciparum.";
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i (Blood fluke).
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                   Han Z.;

"The full-length cDNA sequences of Schistosoma japonicum submitted (NOV-2004) to the EMBL/GenBank/DDBJ databases. EMBL; AY815296; AAW27028.1; -; mRNA.

GO; GO:0005489; F:electron transporter activity; IEA.
GO; GO:0006118; P:electron transport; IEA.
InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin-fold.
Ffam; PP00085; Thioredoxin, 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q5DAX8;
10-MAY-2005 (TrEMBLrel. 30,
10-MAY-2005 (TrEMBLrel. 30,
10-MAY-2005 (TrEMBLrel. 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical protein.
Schistosoma japonicum (Blood fluke).
Eukaryota; Metazoa; Platyhelminthes;
Schistosomatoidea; Schistosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR006662; Thiored.
InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
Pfam; PF00085; Thioredoxin; 1.
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Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; AY673828; ART76629.1; -, mRNA.
GO; GO:0005489; F:electron transporter activity; IEA.
GO; GO:0006118; P:electron transport; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRINTS; PR00421; THIOREDOXIN.
TIGRAMS; TIGRO1068; thioredoxin; 1.
PROSTITE; PS00194; THIOREDOXIN; 1.
SEQUENCE 104 AA; 11995 MW; 8D36E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Beers L.F., Williams
                                                                                                                                                                                                                                                                                                   Hypothetical
SEQUENCE 10
                                                                                                                                                                                                                                                                                                                                               PRINTS; PR00421; THIOREDOXIN.
TIGRFAMS; TIGR01068; thioredoxin; 1.
PROSITE; PS00194; THIOREDOXIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLEOTIDE SEQUENCE.
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SCHJA
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                                                                                                                                                                                                                                                                                                   MW;
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                                                                                                                                                                                                                       Score 47; DB 2; Length 104; Pred. No. 2.5;
                                                                                                                                                                                                                                                                                                   8D36E83058742262 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
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                                                                                                                                                                                                   Mismatches
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databases.
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Q48985;
01-NOV-1996 (TrEMBLrel. 01.
01-NOV-1996 (TrEMBLrel. 01.
01-MAR-2004 (TrEMBLrel. 26.
                                                                                  "Major K.E.;
"Major structural and novel potential virulence mechanisms
genomes of multiple Campylobacter species.";
Submitted (DEC-2004) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an
EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QHIMM5_CAMUP PRELIMINARY;
QHHMM5;
QHHMM5;
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
13-SEP-2005 (TrEMBLrel. 31,
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   SEQUENCE
                                     preliminary data.
EMBL; AAFJ01000012; E
                                                                                                                                                                                                                                                                                                               Fouts D.E., Mongodin E.F., Mandrell R.E., Miller W.G., Rasko D.A.,
Jacques R.J., Brinkac L.M., DeBoy R.T., Parker C.T., Daugherty S.C.,
Dodson R.J., Durkin A.S., Madupu R.R., Sullivan S.A., Shetty J.U.,
Alydeji M.A., Shvartsbeyn A.A., Schatz M.C., Badger J.H., Fraser C.M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Campylobacter upsaliensis RM3195.
Bacteria; Proteobacteria; Epsilonproteobacteria; Campylobacterales; Campylobacteraceae; Campylobacter.
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"Exploring the Mycoplasma capricolum genome: a minimal cell reveals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mycoplasma capricolum.
Bacteria; Firmicutes; Mollicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Thioredoxin.
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STRAIN=ATCC 27343;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=RM3195
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InterPro; IPR005746; Thioredoxin.
InterPro; IPR006663; Thioredox_dom2.
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Mol. Microbiol. 16:955-967(1995).
EMBL; Z33053; CAA83726.1; -; Genomic_DNA.
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   EAL52411.1; -; Genomic DNA.
11271 MW; 3056CAA6990B8F74
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       CRC64;
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)Y9 CLOPA P
Q7M0Y9 CLOPA P
Q7M0Y9;
01-MAR-2004 (7)
01-MAR-2004 (7)
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                                                                                                                          Name=trxA; OrderedLocusNames=BF0642;
Bacteroides fragilis (strain ATCC 25285 / NCTC 9343).
Bacteria; Bacteroidetes; Bacteroidetes (class); Bacteroidales;
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01-FEB-2005 (TrEMBLrel. 29, Last ann
Putative thioredoxin.
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Bacteria; Firmicutes; Clostridia; Clostridiales; Clostridiaceae;
                                                                              Bacteroidaceae; Ba
NCBI_TaxID=272559;
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QSLHJ1;
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InterPro; IPR006663; Thioredox dom2.
InterPro; IPR005746; Thioredoxin.
InterPro; IPR012336; Thioredoxin-like.
InterPro; IPR012335; Thioredoxin fold.
                        NUCLEOTIDE SEQUENCE [LAF
PubMed=15746427; DOI=10.
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TIGRFAMs; TIGR01068; thioredoxin; 1.
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HSSP; P00274; 1F6M.
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                        [LARGE SCALE GENOMIC DNA]
[=10.1126/science.1107008;
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DT 01-UUN-2003 (TrEMBLrel. 24,
DT 01-UUN-2003 (TrEMBLrel. 24,
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DE Thioredoxin.
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OS Bacillus cereus (strain ATC
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Q4L5F0;
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13-SEP-2005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Whole genome sequencing of Staphylococcus haemolyticus uncovers extreme plasticity of its genome and dynamism in the evolution of human-colonizing staphylococcal species.";
Submitted (DEC-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AP006716; BAE05125.1; -; Genomic_DNA.
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Science 307:1463-1465(2005).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Takeuchi F., Watanabe S., Baba T., Yu:
Morimoto Y., Kuroda M., Takahashi M.,
Lee J.C., Hiramatsu K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Staphylococcus haemoly Bacteria; Firmicutes;
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R EMBL; AE017012; AAP11431.1; -; Genomic_DNA.

R HSSP; P80579; INSW.

R GO; GO:0005489; F:electron transporter activity; IEA.

R GO; GO:00054810; P:electron transport; IEA.

R GO; GO:0006810; P:transport; IEA.

R InterPro; IPR006662; Thioredoxin.

R InterPro; IPR005746; Thioredoxin.

R InterPro; IPR005663; Thioredox.

R Pfam; pF00085; Thioredoxin; 1.

R PFATTS; PR00421; THIOREDOXIN.

R PRINTS; PR00421; THIOREDOXIN.

R PROSITE; P800194; THIOREDOXIN; 1.

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O SEQUENCE 104 AA; 11478 MW; B3689CCCC245EE87 CRC64;
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Ivanova N., Sorokin A., Anderson I., Galleron N., Candelon B.,

Kapatral V., Bhattacharyya A., Reznik G., Mikhailova N., Lapidus A.,

Chu L., Mazur M., Goltsman E., Larsen N., D'Souza M., Walunas T.,

Chu L., Mazur M., Goltsman E., Larsen N., D'Souza M., Walunas T.,

Chu L., Pusch G., Haselkorn R., Fonstein M., Ehrlich S.D.,

Chu L., Syrpides N.C.;

Overbeek R., Kyrpides N.C.;

"Genome sequence of Bacillus cereus and comparative analysis with

Bacillus anthracis.";
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Sequence 5, Application (Sequence 5, Applicant) (Sequence 5, Applicant) (Sequence 1, 1985261) (Sequence 1, 198
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Query Match
Best Local Similarity
Matches 6; Conserv
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NAME: Connell, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2879
TELECOMMUNICATION INFORMATION:
TELEPHONE: 303/863-9700
                                                                                                                                                                                                                    IELEFAX: 303/863-0223
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/883,804
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MEDIUM TYPE: Floppy disk
                                                                                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                      TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 514
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US-08-084-718-6
US-08-084-718-8
US-08-084-718-9
US-08-084-718-29
US-08-084-718-29
US-08-084-718-29
US-08-084-718-34
US-08-084-718-34
US-08-084-718-34
US-08-084-718-34
US-08-084-718-34
US-08-084-718-34
US-08-084-718-34
US-08-084-718-36
US-08-084-718-36
US-08-084-718-36
US-08-084-718-36
US-08-084-718-36
US-08-084-718-76-8
US-08-443-976-8
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                                                       DB 1; Length 6;
  Indels
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Sequence 65, Appl
Sequence 18, Appl
Sequence 4, Appl
Sequence 4, Appl
Sequence 4, Appl
Sequence 7037, Ap
Sequence 23, Appl
Sequence 25, Appl
Sequence 27, Appl
Sequence 28, Appl
Sequence 33, Appl
Sequence 34, Appl
Sequence 6, Appli
Sequence 8, Appli
Sequence 8, Appli
Sequence 9, Appli
Sequence 9, Appli
Sequence 5, Appli
Sequence 9, Appli
  0
     Gaps
     0
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FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION ADATA:
APPLICATION NUMBER: JP 5-238402
FILING DATE: 24-SEP-1993
PRIOR APPLICATION NUMBER: PCT/JP94/01572
APPLICATION NUMBER: PCT/JP94/01572
APPLICATION NUMBER: PCT/JP94/01572
APPLICATION NUMBER: PCT/JP94/01572
ATTORNEY/AGENT INFORMATION:
NAME: HALEY Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: SHGN-10
TELECOMMUNICATION INFORMATION:
TELEPAN: (212) 596-9000
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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US-08-627-907A-6
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; TOPOLOGY: not releva:
; MOLECULE TYPE: peptide
US-08-627-907A-6
                                                                                                                                                      US-07-901-713A-3
                                                                                                                                                                        RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application US/08627907A
PATENT NO. 6060302
GENERAL INFORMATION:
APPLICANT: HIRANO, Naoto
APPLICANT: HIRAL, Hisamaru
TITLE OF INVENTION: HUMAN PHOSPHOLIPASE C-ALPHA AND DNA
TITLE OF INVENTION: SEQUENCE ENCODING THE SAME
NUMBER OF SEQUENCES: 8
GENERAL INFORMATION:
APPLICANT: Tuite, Michael F.
APPLICANT: Freedman, Robert B.
APPLICANT: Markus, Henry Z.
APPLICANT: Schultz, Loren D.
APPLICANT: Montgomery, Donna L
                                                                                                                 Sequence 3, Application US/07901713A Patent No. 6291205
                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: c/o FISH & NEAVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 1251 Ave.
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U
                                                                                                                                                                                                                                                                                                                             Local
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6; Conserva
                                                                                                                                                                                                                                  1 WCGPCK 6
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                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                            not relevant
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                                                                                                                                                                                                                                                                                                         100.0%; Score 47; DB 2;
100.0%; Pred. No. 4.6e+05;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                 Length 6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-750-142B-54
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Best Local Similarity
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                                                                                                                                                                                                                                OPERATING SYSTEM: Windows95

OPERATING SYSTEM: Windows95

SOFTWARE: FastSEQ for Windows Version 2.0

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/750,142B

FILING DATE: 25-NOV-1996

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/SE96/00365

FILING DATE: 22-MAR-1996

APPLICATION NUMBER: SE9501067-4

FILING DATE: 24-MAR-1995

ATTORNEY/AGENT INFORMATION:
NAME: PERSET 1895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/07/901,713A
CURRENT FILING DATE: 1992-06-12
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: METHOD FOR INCREASING PRODUCTION OF TITLE OF INVENTION: DISULFIDE BONDED RECOMBINANT PROTEINS BY SACCAROMYCES TITLE OF INVENTION: CEREVISIAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 18469
                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatibl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 6
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                 TELEFAX: 617-542-8906
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                              NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 06:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Sarnstrand, Bengt
TITLE OF INVENTION: NEW PEPTIDES WITH
TITLE OF INVENTION: IMMUNOMODULATORY EFFECTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Bergstrand, Hakan
APPLICANT: Briksson, Tomas
APPLICANT: Lindwall, Magnus
APPLICANT: Sarnstrand, Bengt
                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 225 I
                    TOPOLOGY:
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                                     amino acid
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                                                                                                                      617-542-8906
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                                                                                                                                                                                                                                                                                                                                                                                                                                         IBM Compatible
SYSTEM: Windows95
peptide
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Pred. No. 4.6e+05;
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5210073-1
; PATENT NO. 5210073
; PATENT NO. 5210073
; PATENT YODOI, JUNJI; UCHIDA, ATSUSHI; TAGAYA, YUTAKA
; MITSUI, AKKRA; HIRAKAWA, TADASHI
; TITLE OF INVENTION: METHOD FOR TRREATING CANCER THERAPY
; RADIATION DAMAGE OR ARTERIOSCLEROSIS USING HUMAN ADF
; NUMBER OF SEQUENCES: 1
; CURRENT APPLICATION NUMBER: US/07/589,616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 6
US-08-826-910-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 28-SEP-1990
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                          OPERATING SYSTEM: DOS
SOFTWARE: FRATSEQ for Windows Version 2.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/826,910
FILING DATE: Herewith
CLASSIFICATION 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Shah, Purvi
                                                      NAME: Billings, Lucy J.

NAME: Billings, Lucy J.

REGISTRATION NUMBER: 97-0

REJECOMPUNICATION INFORMATION:

TELEPHONE: 415-855-0555

TELEPAX: 415-845-4166
                                INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 105 amino acids
                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                           ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                          USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Incyte Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            THIOREDOXIN-LIKE PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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                                                                                                               PF-0268 US
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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Sequence 4, Application US/08826910

Patent No. 5919657

GENERAL INFORMATION:
APPLICANT: Hilman, Jennifer L.
APPLICANT: Shah, Purvi
TITLE OF INVENTION: THIOREDOXIN-LIKE PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
В
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                                                                                                             US-08-826-910-4
                                                        Matches
                                                                      Best Local Similarity
                                                                                 Query Match
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                                                                                                                                                                                                TELEX:
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 105 mnino acids
                                                                                                                                                                                                                                                                            CLASSIFICATION: 530
PRIOR APPLICATION NUMBER:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/826,910
FILING DATE: Herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                    IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 3174 Por
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LIBRARY: GenBa
                                                                                                                          LIBRARY: GenBank
CLONE: 453972
                                                                                                                                                                    STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                             TYPE: amino acid
                                                                                                                                                                                                                                                                   TELEFAX: 415-845-4166
 31
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                           WCGPCK 6
 WCGPCK 36
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                                                       Conservative
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                                                                    100.0%;
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                                                                   Score 47; DB 1; Length 105; Pred. No. 4.4;
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                                                       Indels
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                                                       Gaps
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RESULT 8

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Query Match
Best Local Similarity
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SOFTWARE: PATENTIN Ver. 2.
SEQ ID NO 68
LENGTH: 105
TYPE: PRT
                                                                                                                              RESULT 10
US-08-386-729A-10
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                                                                                                                                                                                                                                                                                                                                                 ; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-513-999C-8037
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US-09-601-144-68
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                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins. 6783961
FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 8037
LENGTH: 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Wright, Jim A.
APPLICANT: Young, Aiping H.
APPLICANT: Young, Aiping H.
APPLICANT: Lee, Yoon S.
TITLE OF INVENTION: OLIGONUCLEOTIDE SEQUENCES COMPLEMENTARY TO THIOREDOXIN
TITLE OF INVENTION: AND THIOREDOXIN REDUCTASE GENES AND METHODS OF USING
TITLE OF INVENTION: SAME TO MODULATE CELL GROWTH
FILE REFERENCE: 683-112US-A
CURRENT APPLICATION NUMBER: US/09/601,144
CURRENT FILING DATE: 2000-10-18
CURRENT FILING DATE: 2000-10-18
                                                                                    Sequence 10, Application US/08386729A
Patent No. 5753435
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Sequence 68, Appro-
Sequence 68, Appro-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/073,196
PRIOR FILING DATE: 1998-01-30
                                                                         GENERAL INFORMATION:
                                   APPLICANT:
 APPLICANT:
APPLICANT:
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Aharonowitz, Yair
Van Der Voort, Lucia H.
Cohen, Gerald
Bovenberg, Roelof A. L.
                                                                                                                                                                                                                                                                                             100.0%; Score 47; DB 100.0%; Pred. No. 4.4;
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RESULT 11
US-07-745-382-22
; Sequence 22, Application US/07745382
------ No. 5270181
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Best Local Similarity
Thes 6; Conserve
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APPLICATION NUMBER: US 08/274,043
FILING DATE: 12-JULY-1994
APPLICATION NUMBER: US 07/820,688
FILING DATE: 24-MARCH-1992
APPLICATION NUMBER: PCT/NL91/000101
FILING DATE: 18-JUNE-1991
APPLICATION NUMBER: EP 90201598.1
PILING DATE: 18-JUNE-1990
ATTORNEY/AGENT INFORMATION:
NAME: DBSGLUGGERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
              APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Like Molecules
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Rae-Venter, Barbara
REGISTRATION NUMBER: 32,750
REFERENCE/DOCKET NUMBER: GBRO-024/02US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 926-6200
TELEFAX: (415) 854-3713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/386,729A
FILING DATE: 10-FEB-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Kattevilder, Alfred
APPLICANT: Pallissa, Harriet
TITLE OP INVENTION: An oxido reductase enzyme system
TITLE OP INVENTION: obtainable from P. chrysogenum, the set of genes
TITLE OF INVENTION: encoding the same and the use of oxido reductase enzyme
TITLE OF INVENTION: systems or genes encoding the same for increasing antibiotic
TITLE OF INVENTION: production
NUMBER OF SEQUENCES: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A. ZIP: 94025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
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    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                   31 WCGPCK 36
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                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 47; DB 1; Length 106; ilarity 100.0%; Pred. No. 4.5; Conservative 0; Mismatches 0; Indels
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Kattevilder, Alfred
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Genetics Institute, Inc.
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTMARE: PATENTIA Release #1.0, Version #1.25
SOPTMARE: DATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/745,382
FILING DATE: 1991.0814
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/652,531
FILING DATE: 06-FEB-1991
ATTORNEY/AGENT INFORMATION:
NEMBER: CEST. TIMED
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 5292646
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (617) 876-585
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
            FILING DATE: 19920728
CLASSIFICATION: 435
PRIOR APPLICATION DATA: 07/
APPLICATION NUMBER: US 07/
FILING DATE: 06-FEB-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
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CITY: Car
STATE: M:
COUNTRY:
                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/921,848
FILING DATE: 19920728
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                  COUNTRY: U.ZIP: 02140
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FILING DATE:
                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
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 14-AUG-1991
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-- NO: 22:
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                                                                 US 07/652,531
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Pred. No. 4.6;
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ATTORNEY/AGENT INFORMATION:

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; LENGTH: 109 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-165-301A-22
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                 Query Match
Best Local Similarity
Watches 6; Conserve
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US-08-165-301A-22
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                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI 5
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-1551
INFORMATION FOR SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FOR COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/165,301A
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Grant, Kathleen
APPLICANT: LAVALIIE, Edward R.
APPLICANT: LAVALIIE, EDWARD R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES,
THIORE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: McCoy, John
APPLICANT: Diblasio-Smith, Elizabeth
                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 109 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE
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T: U.S.A.
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87 CambridgePark Drive
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Pred. No. 4.6;
                                                Score 47; DB 1; Length 109; Pred. No. 4.6;
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US-08-165-301A-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-165-301A-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC COMPATIBLE COMPATION SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, Ve CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/165,301A FILING DATE: CLASSIFICATION: 435
AFTORNEY/AGENT INFORMATION: 435
ANAME: Meinert, M. C. REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: G1 5188D TELEFDANE: (617) 876-1170
TELEFPANE: (617) 876-1170
TELEFPANE: (617) 876-5851
INFORMATION FOR SEQ ID NO: 26:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2
Patent No.
                                                                                                                                                                                                                            Sequence 28, Appli
Patent No. 5646016
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                          GENERAL INFORMATION:
                               APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS
LENGTH: 109 amino aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 2:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: McCoy, John APPLICANT: DiBlasio-Sm
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 STREET:
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                     ADDRESSEE:
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5. 5646016
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1: U.S.A.
                                                                                                                                                                                                                                                    Application US/08165301A
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B: Genetics Institute,
87 CambridgePark Drive
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LaVallie, Edwar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DiBlasio-Smith,
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COUNTRY:

Massachusetts: U.S.A.

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Best Local Similarity
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            COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/810,436
FILING DATE: 04-MAR-1997
CLASSIFICATION + 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/165,301
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/165,301A
FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: G1 5188D
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPHONE: (617) 876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 6143524
GENERAL INFORMATION:
APPLICANT: MCCOy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 22,
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INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: DiBlasio-Smith, Elizabeth
APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: 1i
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                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Cambridge
STATE: Massing
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REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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GI 5188D
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RESULT 17
US-08-810-436-26
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                                                                   US-08-810-436-26
                                                                                                                                                                                                                                           ZIP: 02140

ZIP: 02140

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/810,436

FILING DATE: 04-MAR-1997

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/165,301

FILING DATE: 10-DEC-1993

ATTORNEY/AGENT INFORMATION:

NAMB: Meinert, M. C.

REGISTRATION NUMBER: 33,544

PRIOR APPLICATION NUMBER: 33,544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ":-TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 22
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acids
TYPE: amino acid
TOPOLOGY: linear
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 26, Application US/08810436 Patent No. 6143524
                                Query Match
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APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith,
                                                                                                                                TELEFAX: (617) 876-585
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: T
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: protein
                                                                                MOLECULE TYPE: protein
                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                                                                                                   TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
Local Similarity 100.0%;
es 6; Conservation
                                                                                                                                                                                                     TELEPHONE:
                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Cambridge
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87 CambridgePark Drive
                                                                                                                                  109 amino acids
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   Conservative
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LaVallie, Edwar
                                                                                                                                                                  876-5851
NO: 26:
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-- NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , Edward R.

PEPTIDE AND PROTEIN FUSIONS TO
THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
                 100.0%; Score 47; DB 2; 100.0%; Pred. No. 4.6;
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   Mismatches
                                 Length 109;
   Indels
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Sequence 17, Application US/09485885
PATENT NO. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Cerard, Catherine Marie Eva Fernande
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
                                                                                                                                                                                                                         RESULT 19
US-09-485-885-17
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US-08-810-436-28
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve-
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/810,436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
                                     FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (617) 876-1
TELEFAX: (617) 876-585
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/165,301
FILING DATE: 10-DEC-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith, Elizabeth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0 FILING DATE: 04-MAR-1997 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
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6143524
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
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87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                109 amino acids
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(617) 870: 28:
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                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                       B45107
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                                                                                                                                                                                                                                                      US-09-248-796A-19911
                                                                                                                                                                                                                                                                       RESULT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: amino acid of thioredoxin US-09-166-966E-10
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     GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: WUCLEIC ACID AND AMINO ACID SEQUENCES RELATING:
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT PILING DATE: 1999-02-13
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1998-08-13
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LENGTH: 109
TYPE: PRT
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APPLICANT: CHEN, DAVID CHANHAN
APPLICANT: HU, WIEN-TAI
APPLICANT: CHEN, YUN-JU
APPLICANT: HSEU, TZONG-HSIUNG
APPLICANT: HSEU, TZONG-HSIUNG
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SEQ ID NO 10
LENGTH: 109
TYPE: PRT
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PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB:
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: TW 86114750 PRIOR FILING DATE: 1997-10-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: CHIMERIC PROTEINS WITH A CELLULOSE BINDING FILE REFERENCE: 32350-150960 CURRENT APPLICATION NUMBER: US/09/166,966E CURRENT FILING DATE: 1998-10-06
NUMBER OF SEQ ID NOS: 28208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: thioredoxin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: 1..109
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Pred. No. 4.6
0; Mismatches
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Pred. No.
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SEQ ID NO 19911

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RESULT 23
PCT-US94-14179-26
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PCT-US94-14179-22
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Best Local Similarity
Watches 6; Conserva
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                                               Sequence 26, Application PC/TUS9414179 GENERAL INFORMATION:
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Best Local Similarity
Matches 6; Conserv
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: GI S.
REFERENCE/DOCKET NUMBER: GI S.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPHAX: (617) 876-5851
TELEFAX: 700 ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Candida albicans-
-09-248-796A-19911
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION:
ATTORNEY AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14179
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                          ENGTH:
                                                                                                                                                32
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McCoy, John
DiBlasio-Smith, Elizabeth
Grant, Kathleen
                                                                                                                                                                                                                                                                                                                                            109 amino acids
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Pred. No. 4.6;
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Pred. No.
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RESULT 24
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CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI 7
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acide
TYPE: amino acide
TYPE: amino acide
TYPE: amino acide
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GENERAL INFORMATION:
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TITLE OF INVENTION: PEP-
TITLE OF INVENTION: THIO
TITLE OF INVENTION: THIO
TITLE OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/14179
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Grant, Kathleen
APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: protein
                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                          ZIP: 02140
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics
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                                                                                                                                                                                                                               STREET: 87 Cambi
CITY: Cambridge
STATE: Massachus
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                    APPLICATION NUMBER: FILING DATE:
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ZIP: 02140
CLASSIFICATION:
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87 CambridgePark Drive
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VENTION: PEPTIDE AND PROTEIN FUSIONS TO

VENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES,

VENTION: THIOREDOXIN-LIKE MOLECULES
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                                                                                                                                            Floppy disk
                                        PCT/US94/14179
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US-09-134-001C-3859
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; OTHER INFORMATION: Description of Artificial Sequence: consensus
; OTHER INFORMATION: sequence
US-09-264-419C-6
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US-09-264-419C-6
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          Sequence 3859, Application US/09134001C

Patent No. 6380370

GENERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS

TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: GTC-007

CURRENT APPLICATION NUMBER: US/09/134,001C

CURRENT FILLING DATE: 1998-08-13

PRIOR APPLICATION NUMBER: US 60/064,964

PRIOR PILLING DATE: 1997-11-08

PRIOR APPLICATION NUMBER: US 60/055,779

PRIOR APPLICATION NUMBER: US 60/055,779

PRIOR PILLING DATE: 1997-08-14

NUMBER OF SEQ ID NOS: 5674

SEQ ID NO 3859
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SEQ ID NO 6
LENGTH: 111
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Best Local S
Matches 6
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Patent No. 6174682
GENERAL INFORMATION:
APPLICANT: Khodadoust, Mehran
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: No. 6174682el Thioredoxin Family Active Site Molecules and Uses TITLE OF INVENTION: Therefor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 28:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 109 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 109 amino acids
TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                         {\bf 3}
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US-09-543-681A-4397
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                                                                                                                                 ; ORGANISM: Proteus mirabilis US-09-543-681A-4397
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US-09-882-835-5
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                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: GARY BRETON
APPLICANT: GRAY BRETON
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABIT
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.1002-001
CURRENT APPLICATION NUMBER: US/09/543,681A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR FILING DATE: 1999-04-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 6462187
GENERAL INFORMATION:
APPLICANT: Bandaru,
                                                                                                                                                                 NUMBER OF SEQ ID NOS: 8344
SEQ ID NO 4397
LENGTH: 122
TYPE: PRT
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Best Local Similarity
Matches 6; Conser
                                                                                Query Match
Best Local (
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                                                                 Matches
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CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/211,673
PRIOR FILING DATE: 2000-06-15
NUMBER OF SEQ ID NOS: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bandaru, Rajasekhar
TITLE OF INVENTION: 22109, A NOVEL HUMAN THIOREDOXIN FAMILY
TITLE OF INVENTION: MEMBER AND USES THEREOF
FILE REFERENCE: 10448-063001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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 46 WCGPCK 51
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6; Conserv
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                                                            100.0%; ilarity 100.0%; Conservative 0;
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100.0%; Pred. No.
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                                                                Score 47; DB
Pred. No. 5;
); Mismatches
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RESULT 31
US-09-272-342B-5
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Best Local Similarity
Matches 6; Conserve
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US-09-489-039A-8017
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISSASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
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CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 8017
                                                                                                                                    Sequence 5, Application US/09272342B Patent No. 6294341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 9663
LENGTH: 125
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                                                                                                                     GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Human
-09-949-016-9663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence 9663, Application US/09949016
Patent No. 6812339
                                                                          APPLICANT:
                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2000-09-08
                                      TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
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TYPE: PRT
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KIM, SUNG-HOU
RYU, JAE-RYEON
VENTION: METHOD FOR DETECTING A SUBSTANCE HAVING AN ACTIVITY TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gary Breton et. al
                                                                                             YU, YEON-GYU
                                                                                                                                                                                                                                                                                                                                          100.0%; Score 47; DB 2; Length 125; ilarity 100.0%; Pred. No. 5.1; Conservative 0; Mismatches 0; Indels
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Pred. No. 5.
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US-08-961-083-40
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; LENGTH: 134
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-272-342B-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 40, Application US/08961083 Patent No. 6159469
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                                 Matches
                                                              Query Match
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                          TELEPAX: (301)
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE HARACTERISTICS:
SEQUENCE HARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/272,342B
CURRENT FILING DATE: 1999-03-19
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
FILE_REFERENCE: 290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE:
                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: BROOKES, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                             MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: HP VECU
                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Rockville
STATE: Maryland
                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 20850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Human Genome Sciences, STREET: 9410 Key West Avenue
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Local Similarity 100.0%;
les 6; Conservative 0
                                                Local
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                               Similarity 6; Conser
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 WCGPCK 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Choi et.
                               100.0%; Score 47; DB .larity 100.0%; Pred. No. 5.5; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HP Vectra 486/33
                                                                                                                          linear
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ION: PROTEIN USED FOR SAID METHOD
2901-0125-0
                                                                                                                                                                                                                                       (301) 309-8504
                                                                                                                                         single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Streptococcus pneumoniae Antigens and Vaccines 452
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.50 inch, 1.4Mb storage
                                                                                                                                                                                                                                                                       PB340P2
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                                                             DB 2; Length 137;
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                               Gaps
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TOPOLOGY: linear;
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 40:
US-09-536-784-40
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US-09-765-271-40
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US-09-536-784-40
                                                                                                                                                                                                      PATENT NO. 6887663

PATENT INFORMATION:

APPLICANT: Choi et. al.

TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
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Patent No. 6573082
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                Sequence 40, Application US/09765271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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INFORMATION FOR SEQ ID NO: 40:
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FILING DATE: OCT-30-1997
ATTORNEY/AGENT INFORMATION:
NAME: MICHELLE S. MARKE
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/536,784
FILING DATE: 30-Oct-1997
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Choi et. al. TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines NUMBER OF SEQUENCES: 452
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
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ADDRESSEE: Human Genome Sciences, Inc.
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                                                                               ZIP:
                                                                                                                     CITY: Rockville
STATE: Maryland
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                                                                                                    COUNTRY: USA
                                                                                                                                                                STREET: 9410 Key West Avenue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 137 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Human Genome Sciences, Inc
OPERATING SYSTEM: MSDOS version 6.2
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APPLICATION NUMBER: 09/536,784

FILING DATE: CUNKNOWN
APPLICATION NUMBER: 08/961,083

PILING DATE: COT-30-1997

ATTORNEY/AGENT INFORMATION:

NAME: Michelle S. Marks
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3

TELEPHONE: (301) 309-8504

TELEPAX: (301) 309-8502

INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTER:5TICS:
SEQUENCE CHARACTER:5TICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 40, Application Us
Patent No. 6929930
GENERAL INFORMATION:
APPLICANT: Choi et.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/961,083
FILING DATE: OCT.30-1997
ATTORNEY/AGENT INFORMATION:
NAME: Lin J. Hymel
REGISTRATION NUMBER: 45,414
REFERENCE/DOCKET NUMBER: PB340P2C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: Dell Latitude C610
OPERATING SYSTEM: Windows 2000
SOFTWARE: ASCII Text
                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/765,272A
FILING DATE: 22-Jan-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO: 40:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Streptococcus pneumoniae Antigens and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 WCGPCK 66
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 137 amino acids
TYPE: amino acid
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                                     TELEPHONE: (301)
TELEFAX: (301) 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Maryland
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 20850
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100.0%; Pred. No. 5.5;
tive 0; Mismatches
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                                       309-8439
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                                                               610-5790
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US-09-328-352-5360
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                                                                          ; LOCATION: 1..109
; OTHER INFORMATION: Combined amino acid of thioredoxin and an artificial sequence
US-09-166-966E-8
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// ORGANISM: Acinetobacter baumannii
US-09-328-352-5360

                                                                                                                                                                                           APPLICANT: CHEN, YUN-JU
APPLICANT: HSEU, TZONG-HSIUNG
APPLICANT: HSEU, TZONG-HSIUNG
TITLE OF INVENTION: CHEMERIC PROTEINS WITH A CELLULOSE BINDING DOMAIN
FILE REFERENCE: 32350-150960
CURRENT FILING DATE: 1998-10-06
CURRENT FILING DATE: 1998-10-06
PRIOR APPLICATION NUMBER: TW 86114750
PRIOR FILING DATE: 1997-10-08
NUMBER OF SEQ ID NOS: 11
SEQ ID NO 8
LENGTH: 159
TYPET. DET
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Matches
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SEQ ID NO 5360
LENGTH: 145
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: CHEN, DAVID CHANHAN
APPLICANT: HU, NIEN-TAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: GATY L. Breton et al.
APPLICANT: INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5360, Application US/09328352 Patent No. 6562958
Query Match
Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local Similarity
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                                                                                                                        PEATURE:
NAME/KEY: thioredoxin
                                                                                                                                                              TYPE: PRT
ORGANISM: Unknown
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     100.0%; ilarity 100.0%; Conservative 0
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     Score 47; DB 2; Length 159; Pred. No. 6.3; o; Mismatches 0; Indels
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RESULT 39
US-08-775-978-3
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US-08-775-978-1
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                                                                                                                                                                        Sequence 3, Application US/08775978 Patent No. 5831049
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                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                              GENERAL INFORMATION:
APPLICANT: HILLMAN, JENNIFER L.
APPLICANT: GOLI, SURYA K.
TITLE OF INVENTION: NOVEL HUMAN THIOREDOXIN
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: HILLMAN, JENNIFER L.
APPLICANT: GOLI, SURYA K.
TITLE OF INVENTION: NOVEL HUMAN THIOREDOXIN
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/775,978
FILING DATE: TO BE Assigned
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                   IMMEDIATE SOURCE:
CLONE: 1430906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PROPERTY NAME OF THE PROPERTY NAME OF T
             STREET: 3174 POI
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 3174 POI
CITY: Palo Alto
STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE:
                            ADDRESSEE: Incyte Pharmaceuticals, Inc
STREET: 3174 Porter Drive
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Gaps

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FILE REFERENCE: 2022US
CURRENT APPLICATION NUMBER: US/09/741,243C
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: 60/171,785
PRIOR FILING DATE: 1999-12-22
NUMBER OF SEQ ID NOS: 8
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 167
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US-09-741-243C-4
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APPLICANT: Crawford Jr., John Milton
APPLICANT: Rice, John
APPLICANT: Sevala, Veeresh
APPLICANT: Stewart, Sandy
TITLE OF INVENTION: NOVEL PLANT PORPHOBILINOGEN AND FUSION
TITLE OF INVENTION: PROTEIN THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/09741243C Patent No. 6399352
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                                                           Best Local Similarity
Matches 6; Conserv
                                                                                          Query Match
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 166 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                        ORGANISM: Unknown FEATURE:
                                                                                                                                                                                       TYPE: PRT
                                                                                                                                         OTHER INFORMATION: Thioredoxin functional fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/775,978 FILING DATE: To Be Assigned PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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IMMEDIATE SOURCE:
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CLONE: 1545817
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 WCGPCK 37
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                                                                            Score 47; DB 2; Length 167; Pred. No. 6.5;
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Query Match
Best Local Similarity
Marches 6; Conserv
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US-09-270-767-47690
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US-09-270-767-32473
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 47690
LENGTH: 170
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NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PATENTIN VEr. 2.0
SEQ ID NO 32473
LENGTH: 170
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Patent No. 6703491
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7336-094
CURRENT APPLICATION NUMBER: US/09/270,767
                                                                                     GENERAL INFORMATION:
APPLICANT: YU, YEON-GYU
APPLICANT: KIM, SUNG-HOU
APPLICANT: RYU, JAE-RYEON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 47690, Application US/09270767 Patent No. 6703491
                                                                                                                                                          Sequence 6, Application US/09272342B Patent No. 6294341
                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
              TITLE OF INVENTION: METHOD FOR DETECTING A SUBSTANCE HAVING AN ACTIVITY TITLE OF INVENTION: INHIBIT HIV INFECTION USING IMMUNOASSAY AND VARIANT TITLE OF INVENTION: PROTEIN USED FOR SAID METHOD FILE REFERENCE: 2901-0125-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT ORGANISM: Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Drosophila melanogaster
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   APPLICATION
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NUMBER: US/09/272,342B
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100.0%; Pred. No. 6.
tive 0; Mismatches
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Pred. No.
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Best Local Similarity
Thes 6; Conserv
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                                                                                                                                                                                                  Sequence 5104, Application.

Sequence 500744

Patent No. 6800744

GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
APPLICANT: Lynn A Doucette-Stamm and David Bush
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/583,110
CURRENT FILING DATE: 200-05-26
PRIOR APPLICATION NUMBER: US 09/107,433
PRIOR FILING DATE: 1998-06-30
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR FILING DATE: 1998-05-12
PRIOR APPLICATION NUMBER: US 60/051,553
PRIOR FILING DATE: 1997-07-02
NUMBER: OF SEQ ID NOS: 5322
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SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating to Streptococcus
TITLE OF INVENTION: Pneumoniae for Diagnostics and Therapeutics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 1999-03-19
NUMBER OF SEQ ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: homo sapiens and HIV hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                     COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                        NUMBER OF SEQUENCES: 52 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5222, Application US/09583110
5. 6699703
                                                                                                                                                                                                                                                                                                                                                                                                                                           80 WCGPCK 85
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COMPUTER: <Unknown>
OPERATING SYSTEM: <Unknown>
                                                                                                                              STREET: 100 Beaver Street CITY: Waltham
                                                                                                              STATE: Massachusetts
                                                                                               COUNTRY: USA
                                                                                                                                                                        ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 47; DB 2; Length 177; ilarity 100.0%; Pred. No. 6.8; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                           02354
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Pred. No. 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 185;
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Query Match

100.0%;

Score 47;

DB 2; Length 209

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-166-966E-11
                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: CHIMERIC PROTEINS WITH A CELLULOSE BINDING DOMAIN FILE REFERENCE: 32350-150960
CURRENT APPLICATION NUMBER: US/09/166,966E
CURRENT FILING DATE: 1998-10-06
PRIOR APPLICATION NUMBER: TW 86114750
PRIOR FILING DATE: 1997-10-08
NUMBER OF SEQ ID NOS: 11
SEQ ID NO 11
LENGTH: 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 11, Application US/09166966E Patent No. 6407208
GENERAL INFORMATION:
                                                                                        FEATURE:
NAME/KEY: thioredoxin
LOCATION: 1..109
NAME/KEY: Cellose-binding domain
LOCATION: 162..197
NAME/KEY: RGD
LOCATION: 204..206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: CHEN, YUN-JU APPLICANT: HSEU, TZONG-HSIUNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: CHEN, DAVID C
APPLICANT: HU, NIEN-TAI
  OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 5104: SEQUENCE CHARACTERISTICS:
                                               OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/107,433
FILING DATE: 30-Jun-1998
FRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/ 085131
FILING DATE: May 12, 1998
APPLICATION NUMBER: 60/05153
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-011
TELECOMUNICATION INFORMATION:
TELECOMOUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
les 6; Conserv
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LOCATION: (B) LOCATION 1...194
SEQUENCE DESCRIPTION: SEQ ID NO: 5104:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: protein HYPOTHETICAL: YES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 194 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (781)893-5007
TELEFAX: (781)893-8277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 47; DB:
llarity 100.0%; Pred. No. 7.4;
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DAVID CHANHAN
Combined amino acid sequence of thioredoxin, artificial sequence, cellulose-binding domain, and artificial sequence; the two artificial sequences that two artificial sequences flanking the cellulose-binding domain contain restriction sites
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTC-011
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                                                                                                                                                           RESULT 48
PCT-US92-05707-6
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                                                                                                                       Sequence 6, Application PC/TUS9205707 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (312) 474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 210 amino acids
   TITLE OF INVENTION: Hematopoietic Growth Factor Derived from TITLE OF INVENTION: Tlymphocytes and Methods of Use Therefor NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 906866
FILING DATE: 01 July 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
TELEPAX: (312) 474-0448
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Hemat
                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                       Local Similarity tes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Gruber, Lewis S.
REGISTRATION NUMBER: 30,060
REFERENCE/DOCKET NUMBER: 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/01
FILING DATE: 12-JAN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & ADDRESSEE: Borun
 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 60606-6402
                                                                                                                                                                                                                                  106 WCGPCK 111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                   100.0%; Score 47; DB 2; Length 210; ilarity 100.0%; Pred. No. 7.9; Conservative 0; Mismatches 0: Todala
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Choudhury, Chandra
Marshall, O'Toole, Gerstein, Murray & Bicknell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hematopoietic Growth Factor Derived from T Lymphocytes and Methods of Use Therefor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/08/180,371
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; Mismatches
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION NUMBER: US/08/778,717

FILING DATE: 12-DEC-1996

FRIOR APPLICATION 536

PRIOR APPLICATION NUMBER: JP 352225/1995

**TT.ING DATE: 28-DEC-1995

**TT.ING DATE: 28-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
6; Conserva
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US-08-778-717-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; MOLECULE TYPE: protein PCT-US92-05707-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17, App...
8602689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: PCT/US92/05707
FILING DATE: 19920707
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Gruber, Lewis S.
REGISTRATION NUMBER: 30,060
REFERENCE/DOCKET NUMBER: 27620/30933
                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:

APPLICANT: UDBNO, EIICHI
APPLICANT: UDBNOYUKI, FUJII
APPLICANT: OKADA, MASAHISA
TITLE OF INVENTION: FUSED DNA SEQUENCE, FUSED PROTEIN
TITLE OF INVENTION: EXPRESSED FROM SAID FUSED DNA SEQUENCE AND METHOD FOR
TITLE OF INVENTION: EXPRESSING SAID DNA SEQUENCE
NUMBER OF SEQUENCES: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 346-5750
TELEPAX: (312) 984-9740
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: AMINO ACID
TOPOLOGY: line
                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUS'
ADDRESSEE: P.C.
STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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: Illinois
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                                                                                                                                                                                                                                                                            USA
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Pred. No. 7.9;
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REGISTRATION NUMBER:

24,618

2084-031-0

OBLON, NORMAN F.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 5270181
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: McCOy, John
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Like Molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 14,
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Best Local .
                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/745,382
FILING DATE: 19910814
                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/652,531
PILING DATE: 06-FEB-1991
ATTORNEY/AGENT INFORMATION:
NAME: CBerr, Luann
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI5188A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 265 amino acids
                                                                         TELEFAX: (617) 876-5851 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: RECOMBINANT
PUBLICATION INFORMATION:
AUTHORS: NOBUYUKI FUJII ET AL,
AUTHORS: NOBUYUKI FUJII ET AL,
TITLE: FUSED DNA SEQUENCE, FUSED PROTEIN EXPRESSED
TITLE: FROM SALD FUSED DNA SEQUENCE AND METHOD OF
TITLE: EXPRESSING SALD FUSED PROTEIN
RELEVANT RESIDUES IN SEQ ID NO: 17: FROM 1 TO 265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
TELEPHAX: 703-413-2220
                                                        SEQUENCE CHARACTERISTICS:
                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: |
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Mass
COUNTRY: U.
ZIP: 02140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 265 amino acids
TYPE: amino acid
STRANDEDNESS: single
               LENGTH: 296 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     50 WCGPCK 55
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6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genetics Institute, Inc
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RESULT 52
US-08-115-680-4
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Best Local Similarity
Grandhes 6; Conserve
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US-07-921-848-14
                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC COMPACTIBLE
COMPUTER: IBM PC COMPACTIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/921,848
FILLING DATE: 19920728
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/652,531
FILLING DATE: 06-FEB-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/745,382
FILLING DATE: 14-AUG-1991
ATTORNEY/AGENT INFORWATION:
NAME: CGETT, LUADIN
REGISTRATION NUMBER: 31,822
REFERENCE/DOCKET NUMBER: GI5188A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1110
TELEPHONE: (617) 876-1110
                                                                                                                                                     Query Match
Best Local Similarity
Matches 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 5292646
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                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: McCoy, John
APPLICANT: LaVallie, Edward
APPLICANT: LaVallie, Edward
TITLE OF INVENTION: Peptide and Protein Fusions To
TITLE OF INVENTION: Thioredoxin and Thioredoxin-Like Molecules
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
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CITY: Cambridge
STATE: Massachusetts
COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                          LENGTH:
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                                                                              32 WCGPCK 37
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                                                                                                                 WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                 : (617) 876-5851
(617) TD NO: 14:
                                                                                                                                                     100.0%; Score 47; DB 1; Length 296; ilarity 100.0%; Pred. No. 10; Conservative 0; Mismatches 0; Indels
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Sequence 4, Application US/08115680 Patent No. 5437863

GENERAL INFORMATION:

APPLICANT: Williams, David A.

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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/941,372
APPLICATION NUMBER: US 07/941,372
FILING DATE: 02-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Bak, MATY E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: INDIAUSA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 540-9206
TELEPAX: (215) 540-9206
TELEPAX: (215) 540-9218
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
1 ENUTY: 206 Entic 540-818
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/07941372
Patent No. 5460810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Williams, David A.
APPLICANT: Clark, Steven C.
ZIP: 194/,
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION NUMBER: US/07/941,372
                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Method of Treating Cell Damage or TITLE OF INVENTION: Depletion NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/115,680
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TITLE OF INVENTION: Method of Treating Cell Damage or
TITLE OF INVENTION: Depletion
TUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 296 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Howson and Howson
STREET: Spring House Corporate Cntr, P.O. Box 457
CITY: Spring House
                                                                                                                                                                                                       ADDRESSEE: Howson and Howson
STREET: Spring House Corporate Center,
CITY: Spring House
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity les 6; Conserv
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                                                                                                                                                                                          USA
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US-08-165-301A-14
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                                                                          ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-165-301A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 5646016
GENERAL INFORMATION:
Query Match 100.0%; Score 47; DB Best Local Similarity 100.0%; Pred. No. 10; Matches 6; Conservative 0; Mismatches
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Best Local Similarity
                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
TELEPAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: INI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 540-9206
                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 296 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION: NAME: Bak, Mary E.
                                                                                                                               SEQUENCE CHARACTERISTICS LENGTH: 296 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: LaVallie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Grant, Kathleen APPLICANT: LaVallie, Edwar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: McCoy, John APPLICANT: DiBlasio-Smith,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Campring STATE: Massachusetts COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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                                                                                                                           amino acid
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                                                                                                                                                    296 amino acids
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                                     Length 296
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RESULT 55
US-08-810-436-14
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                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                              Sequence 4, Application PC/TUS9308247 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                              APPLICANT: Genetics Institute, Inc., , APPLICANT: 87 CambridgePark Drive, , APPLICANT: Cambridge, MA 02140, USA TITLE OF INVENTION: Method of Treating TITLE OF INVENTION: Depletion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/165,301
FILING DATE: 10-DEC-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE MOLECULES, AND MODIFIED
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
UMMER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc.
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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APPLICANT: LaVallie, Edward R.
                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acide
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 ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DiBlasio-Smith,
                                                                                                                                                                                                                                                                                                                                    100.0%; Score 47; DB ilarity 100.0%; Pred. No. 10; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
Howson and Howson
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04-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          John
                                                                      Method of Treating Cell Damage or
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PRIOR APPLICATION DATA:
APPLICATION UMBER: US 07/941
APPLICATION UMBER: US 07/941
FILING DATE: 02-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: INDI.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 540-9206
TELEPHONE: (215) 540-5818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (215) 540-581
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                             ZIP: 02140

ZIP: 02140

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTION DATA:

APPLICATION NUMBER: PCT/US94/14179

FILING DATE:

FILING DATE:

FILING DATE:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 19477
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Meinert, M. C.
REGISTRATION NUMBER: 33,544
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: LAVAllie, Edward R.
TITLE OF INVENTION: PEPTIDE AND PROTEIN FUSIONS TO
TITLE OF INVENTION: THIOREDOXIN, THIOREDOXIN-LIKE IT
TITLE OF INVENTION: THIOREDOXIN-LIKE MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: McCoy, John
APPLICANT: DiBlasio-Smith,
APPLICANT: Grant, Kathleen
                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 296 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 100.0%; Score 47; DB Local Similarity 100.0%; Pred. No. 10;
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OPERATING SYSTEM: PC-DOS/MS-DOS
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1: U.S.A.
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LaVallie, Edwar
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REFERENCE/DOCKET NUMBER:

GI 5188D

TELECOMMUNICATION INFORMATION:

876-1170

TELEPHONE:

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; NUMBER OF SEQ ID NOS: 1011
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 1011
; LENGTH: 355
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-012-896-1011
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RESULT 59
US-08-683-007A-2
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Best Local Similarity
"hes 6; Conserva
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; MOLECULE TYPE: protein
PCT-US94-14179-14
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INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 296 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/012,896
CURRENT FILING DATE: 2001-12-10
                                                                                                                                                                                                                                                                                                                                APPLICANT: Wantanabe, Yoshihiro
APPLICANT: Meagher, Madeleine Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 210121.427C27
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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Carter, Darrick
Li, Samuel X.
                                                                                                                                                                                                                                                                                                                                                                                               Foy, Teresa
Fanger, Gary R.
                                                                                                                                                                                                                                                                                                                                                                                                                           McNeill, Patricia D.
Houghton, Raymond L.
Vinals de Bassols, Carlota
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Henderson, Robert A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Skeiky, Yasir A.W.
Hepler, William T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Retter, Marc W. Stolk, John A.
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Kalos, Michael D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mitcham, Jennifer L.
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                                                                                                                    Conservative
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                                                                                                                                    100.0%; Score 47; DB 2; Length 355; 100.0%; Pred. No. 12;
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                                                                                                                       Mismatches
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; SEQUENCE CHARACTERISTICS:
; LENGTH: 408 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-683-007A-2
                                                                          ; TYPE: PRT ; ORGANISM: Mus musculus US-09-513-442-2
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                                                                                                                             SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 467
                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/09513442
Patent No. 6387664
Query Match 100.0%; Score 47; DI
Best Local Similarity 100.0%; Pred. No. 15;
Matches 6; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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Best Local S
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APPLICANT: No. 5858724y, Robert E
APPLICANT: Domanico, Michael
APPLICANT: Yaeger, Keith
APPLICANT: Yaeger, Keith
APPLICANT: Kroeker, Warren
TITLE OF INVENTION: Recombinant Rabbit Tissue Factor
NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                       APPLICANT: Ikemoto, Mitsushi
TITLE OF INVENTION: SPARC Fusion Protein and
TITLE OF INVENTION: Same
FILE REFERENCE: HIGAKI-04218
CURRENT APPLICATION NUMBER: US/09/513,442
CURRENT FILING DATE: 2000-02-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 608-251-9166
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Seay, Nicholas J
REGISTRATION NUMBER: 27386
REFERENCE/DOCKET NUMBER: 74
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-251-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 608-251-9166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 16 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/683,007A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Quarles & Brady
STREET: 1 South Pinckney Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                        32 WCGPCK 37
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                     DB 2; Length 467;
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RESULT 61 US-08-260-582-77 밁

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                                                                                                                                                                                                                                                                                                                                        Sequence 77, Application PC/TUS9505471 GENERAL INFORMATION:
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Best Local Similarity
Matches 6; Conserv
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: MCCOy, John M.
APPLICANT: Lu, Zhijian
TITLE OF INVENTION: METHOD OF DETECTING LIGAND
TITLE OF INVENTION: INTERACTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 77:
SEQUENCE CHARACTERISTICS:
LENGTH: 500 amino acids
                                                                                       APPLICATION NUMBER: POINTORMATION FOR SEQ ID NO:
                                                                                                                               TITLE OF INVENTION: METHOD OF DETECTING LIGAND INTERACTIONS
NUMBER OF SEQUENCES: 76
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Meinert, M. C.
REGISTON NUMBER: 31,544
REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08
FILING DATE: 16-JUN-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CORRENT APPLICATION DATA:
CORRENT APPLICATION UNDERS. UT/08/06/56
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CORRESPONDENCE ADDRESS:
                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
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LENGTH: 500 amino acids
TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 275 WCGPCK 280
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87 CambridgePark Drive
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                                                                                                                 PCT/US95/05471
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RESULT 64
US-09-741-243C-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Lyon, Jeffrey A.
APPLICANT: Lyon, Jeffrey A.
APPLICANT: Lyon, Jeffrey A.
APPLICANT: Angov, Evelina
TITLE OF INVENTION: Isolation and Purification of P. falciparum Merozoite
TITLE OF INVENTION: Protein-142 Vaccine
FILE REFERENCE: 003/241/SAP
CURRENT APPLICATION NUMBER: US/10/057,531A
CURRENT FILING DATE: 2002-01-25
PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: US 60/347,564
PRIOR FILING DATE: 2001-10-26
PRIOR FILING DATE: 2001-10-26
                                                                                   NUMBER OF SEQ ID NOS: 8
SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 2
LENGTH: 551
                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Crawford Jr., John Milton
APPLICANT: Rice, John
APPLICANT: Sevala, Veeresh
APPLICANT: Sevala, Veeresh
APPLICANT: Stewart, Sandy
TITLE OF INVENTION: NOVEL PLANT PORPHOBILINOGEN AND FUSION
TITLE OF INVENTION: PROTEIN THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Apple Macintosh Microsoft Word 6.0 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/10057531A Patent No. 6855322
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                       FILE REFERENCE: 2022US
CURRENT APPLICATION NUMBER: US/09/741,243C
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: 60/171,785
PRIOR FILING DATE: 1999-12-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6399352
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial sequence
OTHER INFORMATION: A plant thioredoxin-porphobilinogen
                    ORGANISM: Artificial Sequence FEATURE:
                                                                   TYPE: PRT
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Local Similarity 100.0%;
les 6; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
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Pred. No.
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APPLICANT: Rice, John
APPLICANT: Kloti, Andreas
APPLICANT: Crawford, John
APPLICANT: Crawford, John
APPLICANT: Lanning, Beth
APPLICANT: Lenning, Beth
APPLICANT: Lenning, Beth
ITILE OF INVENTION: METHODS AND COMPOSITIONS FOR THE
ITILE OF INVENTION: INDENTIFICATION OF MODULATORS OF DEOXXXXLULOSE 5-PHOSPHATE
ITILE OF INVENTION: SYNTHASE ACTIVITY
FILE REFERENCE: 2037 US
CURRENT APPLICATION NUMBER: US/09/626,589
CURRENT APPLICATION NUMBER: US/09/626,589
CURRENT APPLICATION NUMBER: US/09/626,589
CURRENT FILING DATE: 2000-07-27
NUMBER OF SEQ ID NOS: 6
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 824
TYPE: PRT
                                    FEATURE:

OTHER INFORMATION: Residues 1-165 are from the thioredoxin sequence OTHER INFORMATION: found in the vector pET32 supplied by No. 6326164ag; OTHER INFORMATION: Residues 166-824 represent the tDXPS sequence from OTHER INFORMATION: Arabidopsis shown in SEQ ID NO:2.

US-09-626-589-3
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CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT FILING DATE: 1999-02-18

PRIOR APPLICATION NUMBER: US 60/074,788

PRIOR PILING DATE: 1998-02-18

PRIOR PILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142
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Best Local S
Matches 6
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Best Local Similarity
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LENGTH: 580
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Query Match
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                       ORGANISM: Artificial Sequence
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100.0%;
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Pred. No. 17;
Score 47;
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DB 2;
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Length 824;
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Query Match
Best Local Similarity
"""hes 6; Conserva
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US-10-237-551-92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Day, Craig H.
APPLICANT: Hosken, Nancy A.
APPLICANT: Hosken, Nancy A.
APPLICANT: Parsons, Joseph M.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AN
TITLE OF INVENTION: TREATMENT OF HERPES SIMPLEX VIRUS INFECTION
FILE REFERENCE: 210121.538C3
CURRENT APPLICATION NUMBER: US/10/237,551
CURRENT FILING DATE: 2002-09-06
NUMBER OF SEQ ID NOS: 254
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 92
LENGTH: 858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 6166178
GENERAL INFORMATION:
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              APPLICATION NUMBER: US/08/
FILING DATE: 19-NOV-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 01-OCT-1996
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                      COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                              APPLICANT: Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                         ZIP: 94111-3834
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cech, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                            CITY: San Francis
STATE: California
                                                                                                                                                                                                                                                                                              ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                              COUNTRY:
APPLICATION NUMBER:
                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
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682151
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                                                                                                                                                                                                                                                                 USA
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Harley, Calvin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lingner, Joachim
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Chapman, Karen B.
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100.0%; Pred. No.
                                                                                                                      US/08/974,549A
US 08/844,419
                                                  08/724,643
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                                                                                                                                                        Version #1.30
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STREET:

Two Embarcadero Center, 8th Floor

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RESULT 69
US-08-912-951-314
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PILING DATE: 01-UC1-L/,
ATTORNEY/AGENT INFORMATION:
AME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1NFORMATION:
TELECOMMUNICATION 1NFORMATION:
                                                                                                                                                                                                                                     Sequence 314, Application US/08912951 Patent No. 6475789
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                  GENERAL INFORMATION:
                                APPLICANT: Nakamura, Toru
APPLICANT: Chapman, Karen B.
APPLICANT: Morin, Gregg B.
APPLICANT: Harley, Calvin
APPLICANT: Harley, Calvin
TITLE OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND
TITLE OF INVENTION: THERAPEUTIC METHODS
NUMBER OF SEQUENCES: 335
                                                                                                                                                                               APPLICANT:
APPLICANT:
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APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/17885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
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APPLICATION NUMBER:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein
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OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino a
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Townsend and Townsend and Crew LLP
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09-MAY-1997
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14-AUG-1997
                                                                                                                                                                                                Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 47;
100.0%; Pred. No.
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enterokinase cleavable, His tagged
thioredoxin moiety and full length hTRT"
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RESULT 70
US-09-402-1818-600
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                                                                                                                                           Sequence 600, Application US/0940
Patent No. 6610839
GENERAL INFORMATION:
GENERAL TAPOLICANT: Cech, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 6; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 09-MAY-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
CLASSIFICATION: 435
CLASSIFICATION: 435
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/912,951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: protein
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LENGTH: 1285 amino aci
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APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-0CT-1996
CLASSIFICATION: 435
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Apple, Randolph REGISTRATION NUMBER: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 18-APR-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 08/854,050
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
           Lingner, Joachim
Nakamura, Toru
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
Harley, Calvin B.
Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
                                                                                                                                                                                                                                                                                                 32 WCGPCK 37
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                                                                                                                                                                                                     Application US/09402181B
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SEQUENCES: 633
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, 576-0300
, 576: 314:
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Pred. No. 35;
                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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NAME/KEY: Protein

LOCATION: 1..1285

OTHER INFORMATION: /note= "fusion protein composed of enterokinase cleavable, His tagged thioredoxin moiety and full length hTRT"

SEQUENCE DESCRIPTION: SEQ ID NO: 600:

US-09-402-181B-600
                                                                                                                    밁
Search completed: February 23, 2006, 00:43:50 Job time: 39.5 secs
                                                                                                                                                                    ફ
                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: AUSENHUB, SCOTT L.
REGISTRATION NUMBER: 42,271
REFERENCE/DOKET NUMBER: 015389-002620US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 600:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WS 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WS 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WS 08/915,503
FILING DATE: 10-OCT-1997
APPLICATION NUMBER: WS 08/915,503
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WS 08/915,503
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WS 08/915,503
FILING DATE: 01-OCT-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/402,181B
FILING DATE: 29-Sep-1997
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                 32 WCGPCK 37
                                                                                                                                                                          1 WCGPCK 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: <Unknown>
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Title:
Perfect score:
Sequence:
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No.
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Maximum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Searched:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein
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   pred. No. score grea and is de:
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80
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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   protein search,
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length: 2000000000
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Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hits satisfying chosen parameters:
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  1867569 seqs,
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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770 771 772 773 774 444 444	64 43 91. 65 43 91. 66 43 91. 67 43 91. 68 43 91.	60 43 91. 61 43 91. 62 43 91. 63 43 91.	56 43 91. 57 43 91. 58 43 91.	53 43 91. 54 43 91. 55 43 91.	50 43 91. 51 43 91. 52 43 91.	47 43 91. 48 43 91. 49 43 91.	43 91.	43 91.	40 43 91. 41 43 91.	37 43 91. 38 43 91.	34 43 91.	31 43 91. 32 43 91.	28 43 91. 29 43 91. 30 43 91.	26 43 91. 27 43 91.	24 43 91. 25 43 91.	22 43 91.	20 43 91.	18 43 91.	16 43 91.	14 43 91.	12 43 91.	10 43 91. 11 43 91.	08 43 91. 09 43 91.	04 43 91. 05 43 91. 06 · · 43 91. 07 · 43 91.
69 43 91.5 74 70 43 91.5 74 71 43 91.5 74 72 43 91.5 74 73 43 91.5 74 74 43 91.5 74 75 43 91.5 74 76 43 91.5 74	64 43 91.5 747 65 43 91.5 747 66 43 91.5 747 67 43 91.5 747 68 43 91.5 747	60 43 91.5 747 61 43 91.5 747 62 43 91.5 747 63 43 91.5 747	56 43 91.5 747 57 43 91.5 747 58 43 91.5 747 50 43 91.5 747	53 43 91.5 747 54 43 91.5 747 55 43 91.5 747	50 43 91.5 747 51 43 91.5 747 52 43 91.5 747	47 43 91.5 747 48 43 91.5 747 49 43 91.5 747	145 43 91.5 747 146 43 91.5 747	143 91.5 747 144 43 91.5 747	40 43 91.5 747 41 43 91.5 747 42 43 91.5 747	30 43 91.5 747 30 43 91.5 747 30 43 91.5 747	33 43 91.5 747 35 43 91.5 747 36 747	31 43 91.5 32 43 91.5	28 43 91.5 747 29 43 91.5 747 30 43 91.5 747	26 43 91.5 747 27 43 91.5 747	24 43 91.5 747 25 43 91.5 747	22 43 91.5 747 23 43 91.5 747	20 43 91.5 74 21 43 91.5 74	11 43 91.5 747 18 43 91.5 747	16 43 91.5 747	14 43 91.5 747	12 43 91.5 747	10 43 91.5 747 11 43 91.5 747	08 43 91.5 747 09 43 91.5 747	04 43 91.5 747 05 43 91.5 747 06 · 43 91.5 747 07 · 43 91.5 747
69 43 91.5 747 70 43 91.5 747 71 43 91.5 747 72 43 91.5 747 73 43 91.5 747 74 43 91.5 747 75 43 91.5 747 76 43 91.5 747	64 43 91.5 747 4 US 65 43 91.5 747 4 US 66 43 91.5 747 4 US 67 43 91.5 747 4 US 68 43 91.5 747 4 US	60 43 91.5 747 4 US 61 43 91.5 747 4 US 62 43 91.5 747 4 US 62 43 91.5 747 4 US 63 43 91.5 747 4 US	56 43 91.5 747 4 US 57 43 91.5 747 4 US 58 43 91.5 747 4 US 50 43 91 5 747 4 US	53 43 91.5 747 4 US 54 43 91.5 747 4 US 55 43 91.5 747 4 US	50 43 91.5 747 4 US 51 43 91.5 747 4 US 52 43 91.5 747 4 US	47 43 91.5 747 4 US 48 43 91.5 747 4 US 49 43 91.5 747 4 US	45 43 91.5 747 4 US	143 91.5 747 4 US	40 43 91.5 747 4 US 41 43 91.5 747 4 US 41 43 91.5 747 4 US	30 43 91.5 747 4 US 37 43 91.5 747 4 US 38 43 91.5 747 4 US	33 43 91.5 747 4 US 34 43 91.5 747 4 US 35 43 91.5 747 4 US	31 43 91.5 747 4 US 32 43 91.5 747 4 US	28 43 91.5 747 4 US 29 43 91.5 747 4 US 30 43 91.5 747 4 US	26 43 91.5 747 4 US 27 43 91.5 747 4 US	24 43 91.5 747 4 US 25 43 91.5 747 4 US	22 43 91.5 747 4 US 23 43 91.5 747 4 US	20 43 91.5 747 4 US	11/ 43 91.5 /4/ 4 US	16 43 91.5 747 4 US	14 43 91.5 747 4 US	12 43 91.5 747 4 US	10 43 91.5 747 4 US 11 43 91.5 747 4 US	08 43 91.5 747 4 US 09 43 91.5 747 4 US	04 43 91.5 747 4 US 05 43 91.5 747 4 US 06 · · 43 91.5 747 4 US 07 · 43 91.5 747 4 US

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Sequence 4, Application US/10304287

| Publication No. US20030083234A1 |
| GENERAL INFORMATION: Waisman, David M. |
| APPLICANT: Waisman, David M. |
| APPLICANT: Kwon, Mijung |
| TITLE OF INVENTION: Anti-angiogenesis methods, compositions and uses therefor |
| FILE REFERENCE: ME02-001 |
| CURRENT APPLICATION NUMBER: US/10/304,287 |
| CURRENT FILING DATE: 2002-11-26 |
| PRIOR APPLICATION NUMBER: US 60/333,866 |
| PRIOR FILING DATE: 2001-11-28 |
| NUMBER OF SEQ ID NOS: 8 |
| SOFTWARE: Microsoft Word |
| LENGTH: 6 |
| TYPE: PRI |
| ORGANISM: mammalian |
                                                                                                    RESULT 2
US-10-660-118A-3
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US-10-304-287-4
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Sequence 3, Application US/10660118A
Publication No. US20040131606A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum
FILE REFERENCE: 2879-98
CURRENT APPLICATION NUMBER: US/10/660,118A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2003-09-10
PRIOR FILING DATE: 2003-04-11
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Best Local Similarity
Matches 6; Conserv
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US-10-170-028-459
US-10-120-028-459
US-10-123-913-426
US-10-017-085A-459
US-10-017-085A-459
US-10-0140-473-426
US-10-140-810-426
US-10-140-816-426
US-10-141-703-426
US-10-141-703-426
US-10-141-703-426
US-10-141-762-426
US-10-141-762-426
US-10-142-428-4-426
US-10-142-428-4-426
US-10-142-428-4-426
US-10-143-027-426
US-10-143-027-426
US-10-143-027-426
US-10-143-027-426
US-10-143-027-426
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US-10-165-038A-459
US-10-165-353A-459
US-10-167-600-459
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Sequence 4, Application US/09801260

Patent No. US20020034801A1

GENERAL INFORMATION:
APPLICANT: Curtis, Rory A.J.
ITILE OF INVENTION: 22105, A NOVEL HUMAN THIOREDOXIN FAMILY
ITILE OF INVENTION: MEMBER AND USES THEREOF
FILE REFERENCE: 10448-022001
CURRENT APPLICATION NUMBER: US/09/801,260
CURRENT FILING DATE: 2001-03-06
PRIOR APPLICATION NUMBER: US 60/187,447
PRIOR FILING DATE: 2000-03-07
NUMBER OF SEQ ID NOS: 5
SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4
LENGTH: 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4, Application US/10735577

Sequence 4, Application No. US20040142897A1

GENERAL INFORMATION:
APPLICANT: Waisman, David M.

TITLE OF INVENTION: Compositions and Methods for Inhibiting Tumuser Application NUMBER: US/10/735,577

CURRENT APPLICATION NUMBER: US/10/735,577

CURRENT FILING DATE: 2003-12-12

PRIOR APPLICATION NUMBER: US 60/433,140

PRIOR APPLICATION NUMBER: US 60/433,140

PRIOR APPLICATION NUMBER: US 60/433,140

SOFTWARE: Microsoft Word

SOFTWARE: Microsoft Word

SOFTWARE: Microsoft Word
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Best Local Similarity
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Best Local Similarity
Watches 6; Conserv?
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US-09-801-260-4
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US-10-735-577-4
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                                                  ; OTHER INFORMATION: consensus sequence US-09-801-260-4
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 3
LENGTH: 6
Best Local Similarity
                 Query Match
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TYPE: PRT
ORGANISM: mammalian
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                                                                                  TYPE: PRT
ORGANISM: Artificial
FEATURE:
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100.0%; Pred. No.
                                                                                                    Sequence
 100.0%;
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 Score 47;
Pred. No.
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 DB 3;
7.1;
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                 Length 52;
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; OTHER INFORMATION: consensus sequence US-10-145-586-43
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APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With

TITLE OF INVENTION: Plants

FILE REFERENCE: 38-21 (5322)B

CURRENT APPLICATION NUMBER: US/10/425,115

CURRENT FILING DATE: 2003-04-28

NUMBER OF SEQ ID NOS: 369326

SEQ ID NO 351019
              Query Match
Best Local S
Matches 6
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CURRENT FILING DATE: 2002-05-14
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 95
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 43
LENGTH: 52
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE: INCOMMENTATION CONTROLLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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                                                                                       OTHER INFORMATION: Clone ID: MRT4577_83299C.1.pep -10-425-115-351019
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APPLICANT: Kapeller-Libermann, Rosana
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS
TITLE OF INVENTION: HUMAN THIOREDOXIN FAMILY MEMBERS, HUMAN LEUCINI
TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Alexandra Glucksmann, Maria
APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: M. Galvin, Katherine
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: La Rosa, Thomas J. APPLICANT: Kovalic, David K.
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                                                                                                                                 LENGTH: 82
TYPE: PRT
ORGANISM: Zea mays
                                                                                                                          FEATURE:
              Local Similarity hes 6; Conserv
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                  Conservative
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                               100.0%; Score 47; DB 4; Length 82; 100.0%; Pred. No. 10;
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US-10-424-599-280571
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-408-765A-312
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US-10-408-765A-312
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                                                                    Matches
                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                               ORGANISM: Glycine max
                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                  FEATURE:
                                                                                       Local
13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30 WCGPCK 35
                                                                  l Similarity
6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Conservative
                                 WCGPCK 6
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APPLICANT: Gibson, Bradford W.

APPLICANT: Taylor, Steven W.

APPLICANT: Glenn, Gary M.

APPLICANT: Glenn, Gary M.

APPLICANT: Warnock, Dale E.

TITLE OF INVENTION: TARGETS FOR THERAPEUTIC INTERVENTION

TITLE OF INVENTION: IDENTIFIED IN THE MITOCHONDRIAL PROTEOME

FILE REFERENCE: 660088.465

CURRENT APPLICATION NUMBER: US/10/408,765A

CURRENT APPLICATION NUMBER: US/10/408,765A

CURRENT FILING DATE: 2003-04-04

NUMBER OF SEQ ID NOS: 3077

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 312
                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement; FILE REFERENCE: 38-21(5323)B; CURRENT APPLICATION NUMBER US/10/424,599; CURRENT FILING DATE: 2003-04-28; NUMBER OF SEQ ID NOS: 285684

SEQ ID NO 280571
LENGTH: 90
TYPE: DDT
                                                                                           ; OTHER INFORMATION: Clone ID: PAT_MRT3847_95379C.1.pep US-10-424-599-280571
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 280571, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ghosh, Soumitra S.
APPLICANT: Fahy, Eoin D.
APPLICANT: Zhang, Bing
APPLICANT: Gibson, Bradford
                                                                                                                                               NAME/KEY: unsure
LOCATION: (1)...(90)
OTHER INFORMATION: unsure at all Xaa locations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/10408765A
o. US20040101874A1
100.0%; Score 47; DB llarity 100.0%; Pred. No. 11; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 47; DB 100.0%; Pred. No. 10;
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                                             DB 4; Length 90;
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  Gaps
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RESULT 11
US-10-032-201B-104
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US-10-032-201B-102
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; ORGANISM: Thiocapsa roseopersicina
US-10-032-201B-143
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US-10-032-201B-143
                                                                                                                                                                                                                                                    ; ORGANISM: Chlamydia muridarum US-10-032-201B-102
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Best Local Similarity luv.
"---hes 6; Conservative
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 102
LENGTH: 102
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 143
LENGTH: 91
                                                                                                                                                               Query Match 100.0%; Score 47; DB 4; Length 102; Best Local Similarity 100.0%; Pred. No. 12; Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 102, Application US/10032201B Publication No. US20030167524A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: MCTHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
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APPLICANT: Deckers, Harm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Van Rooijen, Gijs APPLICANT: Deckers, Harm
                                                                                                                                                                                                                                                                                          TYPE: PRT
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                                                                                    27 WCGPCK 32
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Dalmia, Bipin Kumar
Del Val, Greg
Zaplachinski, Steve
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Heifetz, Peter Bernard
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Heifetz, Peter Bernard
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100.0%; Pred. No. 11;
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GENERAL INFORMATION:

APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Briggs, Steven
APPLICANT: Del Val, Greg
APPLICANT: Del Val, Greg
APPLICANT: Del Val, Greg
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION COMPOSITIONS
FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT APPLICATION OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 105
LENGTH: 102
TYPE: PRT
ORGANISM: Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 12
US-10-032-201B-105
; Sequence 105, App
; Publication No. U
RESULT 13
US-10-022-201B-126
; Sequence 126, Application US/10032201B
; Publication No. US20030167524A1
; GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 104
LENGTH: 102
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Best Local Similarity
Matches 6; Conserv
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Best Local Similarity 100
Matches 6; Conservative
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APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Brigs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
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APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Psittaci
                                                                                                                                                          27 WCGPCK 32
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Zaplachinski, Steve
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                                                                                                                                                                                                                                                Conservative
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100.0%; Pred. No. 12
ative 0; Mismatches
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100.0%; Pred. No. 12;
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RESULT 15
US-10-501-282-3126
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; ORGANISM: Mycoplasma
US-10-032-201B-128
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 128
LENGTH: 102
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 126
LENGTH: 102
                                                   Sequence 3126, Application US/10501282 Publication No. US20050203280A1 GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity
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APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kum
APPLICANT: MCMICHAEL, JOHN CALHOUN
APPLICANT: ZAGURSKY, ROBERT JOHN
APPLICANT: RUSSELL, DAVID PARRISH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS,
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
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                                                                                                                                                                                        29 WCGPCK 34
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                                                                                                                                                                                                                                                               6; Conservative
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Dalmia, Bipin Kumar
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Dalmia, Bipin Kumar
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                                                                                                                                                                                                                                                                                 Score 47;
Pred. No.
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                                                                                                                                                                                                                                                                 Indels
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RESULT 17
US-10-032-585-7785
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US-10-032-201B-96
                                                                                                                                                                                                                                                                                                                                                                                      US-10-032-201B-96
                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 96
LENGTH: 103
TYPE: PRT
                                                                           Sequence 7785, Application US/10032585 Publication No. US20030180953A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 96, Application US/1003220:
Publication No. US20030167524A1
GENERAL INFORMATION:
APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.2
SEQ ID NO 3126
LENGTH: 102
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APPLICANT:
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TITLE OF INVENTION: DOLYPEPTIDE ANTIGENS, IMMUNOGENIC COMPOSITIONS AND USES THEREOF
FILE REFERENCE: AM100780 L2
CURRENT APPLICATION NUMBER: US/10/501,282
CURRENT FILING DATE: 2004-07-09
FRIOR APPLICATION NUMBER: 60/333,777
PRIOR FILING DATE: 2001-11-29
PRIOR FILING DATE: 2002-11-18
PRIOR APPLICATION NUMBER: 60/426,742
PRIOR APPLICATION NUMBER: FCT/US02/36123
FRIOR APPLICATION NUMBER: PCT/US02/36123
FRIOR FILING DATE: 2002-11-25
RUMBER: OF SEQ ID NOS: 6653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
                                                        APPLICANT: Terry, Roemer D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 38814 351B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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Dalmia, Bipin Kumar
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Zaplachinski, Steve
                  Bo, Jiang
Charles, Boone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/10032201B
No. US20030167524A1
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US-10-032-585-7785
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CURRENT FILING DATE: 2001-12-20
RUMBER OF SEQ ID NOS: 8000
SOFTWARE: PatentIn version 3.1
SEQ ID NO 7785
LENGTH: 103
                                                                                                                                                                                                                                                                                           Sequence 270, Application US/10451467A Publication No. US20040161840A1 GENERAL INFORMATION:
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SEQ ID NO 7
LENGTH: 103
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                                                                                                                                                                  APPLICANT: CONTRERAS, ROLAND HENRI
APPLICANT: EBERHARDT, INES
APPLICANT: LUYTEN, WALTER HERMAN MARIA LOUIS
APPLICANT: REEKMANS, RIEKA JOSEPHINA
TITLE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION
TITLE OF INVENTION: YEAST AND FUNGI
                                  CURRENT APPLICATION NUMBER: US/10/451,467A CURRENT FILING DATE: 2003-06-19 PRIOR APPLICATION NUMBER: EP 00870318.3 PRIOR FILING DATE: 2000-12-22 PRIOR APPLICATION NUMBER: EP 01870002.1 PRIOR FILING DATE: 2001-01-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/660,118A CURRENT FILING DATE: 2003-09-10 PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or FILE REFERENCE: 2879-98
PRIOR APPLICATION NUMBER: EP 01870003.9 PRIOR FILING DATE: 2001-01-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Gene Disruption Methodologies for Drug Target Discovery
                                                                                                                                                            FILE REFERENCE: JAB-1667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
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APPLICANT: CONTREAS, ROLAND HENRI
APPLICANT: EBERHARDT, INES
APPLICANT: LUTTEN, WALTER HERMAN MARIA LOUIS
APPLICANT: REEKWANS, RIEKA JOSEPHINA
ITILE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION IN
ITILE OF INVENTION: YEAST AND FUNGI
ITILE OF INVENTION: YEAST AND FUNGI
ITILE OF INVENTION: YEAST AND FUNGI
IFILE REFERENCE: JAB-1667
CURRENT FILING DATE: 2003-66-19
PRIOR APPLICATION NUMBER: US/10/451,467A
CURRENT FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: EP 01870013.3
PRIOR FILING DATE: 2001-01-04
PRIOR FILING DATE: 2001-01-04
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 732
SOFTWARE: Patentin version 3.1
SEQ ID NO 450
LENGTH: 103
TYPE: PRI
ORGANISM: Candida albicans
GENERAL INFORMATION: ROLAND HENRI

APPLICANT: CONTRERAS, ROLAND HENRI

APPLICANT: EBERHARDT, INES

APPLICANT: EBERHARDT, INES

APPLICANT: EBERHARDT, INES

APPLICANT: REEKMANS, RIEKA JOSEPHINA

TITLE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION IN

TITLE OF INVENTION: YEAST AND FUNGI

FILE REFERENCE: JAB-1667

CURRENT APPLICATION NUMBER: US/10/451,467A

CURRENT FILING DATE: 2003-06-19

PRIOR APPLICATION NUMBER: EP 00870318.3

PRIOR APPLICATION NUMBER: EP 01870002.1

PRIOR APPLICATION NUMBER: EP 01870003.9

PRIOR APPLICATION NUMBER: EP 01870003.9
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SEQ ID NO 270
LENGTH: 103
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Best Local Similarity
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100.0%; Pred. No.
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Pred. No. 12;
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NUMBER OF SEQ ID NOS: 732

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                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/240, 498
PRIOR FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: 60/260, 284
PRIOR FILING DATE: 2001-01-08
PRIOR PRIOR PRICING DATE: 2001-01-09
PRIOR PRICING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/264, 794
PRIOR FILING DATE: 2001-01-29
PRIOR PRICING DATE: 2001-01-29
PRIOR PRICING DATE: 2000-10-06
PRIOR FILING DATE: 2000-10-06
PRIOR PRICING DATE: 2000-09-15
PRIOR APPLICATION NUMBER: 60/234, 675
PRIOR APPLICATION NUMBER: 60/237, 675
PRIOR APPLICATION NUMBER: 60/237, 675
PRIOR APPLICATION NUMBER: 60/237, 662
PRIOR APPLICATION NUMBER: 60/237, 662
PRIOR APPLICATION NUMBER: 60/274, 862
PRIOR FILING DATE: 2000-09-15
PRIOR APPLICATION NUMBER: 60/237, 662
PRIOR APPLICATION NUMBER: 60/237, 662
PRIOR PRICING DATE: 2001-03-09
PRIOR PRICING DATE: 2001-03-09
PRIOR PRICING DATE: 2001-03-09
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SEQ ID NO 614
LENGTH: 103
TYPE: PRT
ORGANISM: Candida albicans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 62, Application US/09954342 Publication No. US20030170838A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 6; Conserv
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APPLICANT:
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APPLICANT: SPYTEK, KIMBERLY ANN
APPLICANT: TAUPIER, RAYMOND J.
APPLICANT: VERNET, CORINE A.
APPLICANT: COLMAN, STEVEN D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/954,342
CURRENT FILING DATE: 2001-09-17
CURRENT FILING UNUMBER: 60/233,382
PRIOR APPLICATION NUMBER: 60/233,382
PRIOR FILING DATE: 2000-09-18
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APPLICANT: ZERHUSEN, BRYAN
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AND POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 21402-114
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                                       APPLICATION NUMBER: 60/
                                                                                                         APPLICATION NUMBER: 60/232,676
                                                                                                                                                   APPLICATION NUMBER: 60/274,862
FILING DATE: 2001-03-09
APPLICATION NUMBER: 60/233,801
                                                                                      FILING DATE:
                FILING DATE: 2000-09-20
APPLICATION NUMBER: 60/233,402
                                                                                                                                   FILING DATE:
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GERLACH, VALERIE I.
MACDOUGALL, JOHN R.
MACDOUGALL, JOHN R.
PRIJER
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GUNTHER, ERIK
ELLERMAN, KAREN
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                                                                                      2000-09-1
                                                                                                                                   2000-09-19
2000-09-
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                                                               60/233,960
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Pred. No.
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APPLICANT: EASTELLI, LUCA
APPLICANT: ZERHUSEN, BRYAN
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY
FILE REFERENCE: 21402-114
CURRENT FILING DATE: 2001-09-17
CURRENT FILING DATE: 2001-09-17
PRIOR APPLICATION NUMBER: 60/233,382
PRIOR APPLICATION NUMBER: 60/240,498
PRIOR FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: 60/240,284
PRIOR APPLICATION NUMBER: 60/260,284
PRIOR FILING DATE: 2001-01-08
PRIOR FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 60/260,973
PRIOR APPLICATION NUMBER: 60/264,794
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/264,794
PRIOR FILING DATE: 2001-01-29
PRIOR APPLICATION NUMBER: 60/238,398
PRIOR FILING DATE: 2000-10-61
PRIOR APPLICATION NUMBER: 60/332,675
PRIOR APPLICATION NUMBER: 60/332,675
PRIOR PILING DATE: 2000-09-15
PRIOR PILING DATE: 2000-09-15
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PRIOR FILING DATE: 2000-09-19
; PRIOR, APPLICATION NUMBER: 60/232,679
PRIOR FILING DATE: 2000-09-15
; NUMBER OF SEQ ID NOS: 104
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 62
; LENGTH: 104
; TYPE: PRT
; ORGANISM: Ovis aries
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US-09-954-342-63
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publication No. US2
; GENERAL INFORMATION;
; APPLICANT: MISHRA,
; APPLICANT: SPYTEK,
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Best Local S
Matches 6
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PRIOR FILING DATE: 2001-03-09
PRIOR APPLICATION NUMBER: 60/233,801
PRIOR FILING DATE: 2000-09-19
PRIOR APPLICATION NUMBER: 60/232,676
PRIOR FILING DATE: 2000-09-15
PRIOR APPLICATION NUMBER: 60/233,960
PRIOR APPLICATION NUMBER: 60/233,960
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GERLACH, VALERIE L.
GERLACH, JOHN R.
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ELLERMAN, KAREN
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STONE, DAVID
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MACDOUGALL, UCLE.

MACDOUGALL, UCLE.
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10. US20030170838A1
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; TYPE: PRT
; ORGANISM: BOS taurus
US-09-954-342-63
                                                    PRIOR APPLICATION NUMBER: 60/233,382
PRIOR FILING DATE: 2000-09-18
PRIOR PELICATION NUMBER: 60/240,498
PRIOR FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: 60/260,284
PRIOR FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 60/260,973
PRIOR PILING DATE: 2001-01-11
PRIOR PILING DATE: 2001-01-11
PRIOR PILING DATE: 2001-01-29
PRIOR PILING DATE: 2001-01-29
PRIOR PILING DATE: 2000-10-06
PRIOR PILING DATE: 2000-10-06
PRIOR PILING DATE: 2000-10-06
PRIOR PILING DATE: 2000-10-06
PRIOR PILING DATE: 2000-09-15
PRIOR FILING DATE: 2000-09-15
PRIOR FILING DATE: 2000-09-15
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Matches
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 63
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APPLICANT:
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APPLICANT: SPYTEK, KIMBERLY ANN
APPLICANT: TAUPIER, RAYMOND J.
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                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY FILE REFERENCE: 21402-114
CURRENT APPLICATION NUMBER: US/09/954,342
CURRENT FILING DATE: 2001-09-17
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FILING DATE: 2000-09-19
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                  APPLICATION NUMBER: 60/274,862 FILING DATE: 2001-03-09
APPLICATION NUMBER: 60/233,801
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6; Conserva
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PEYMAN, JOHN
STONE, DAVID
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RASTELLI, LUCA
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GERLACH, VALERIE L.
MACDOUGALL, JOHN R.
SMITHSON, GLENNDA
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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APPLICANT: LI, LI
APPLICANT: RASTELLI, LUCA
APPLICANT: RESTRUSEN, BRYAN
TITLE OF INVENTION: NOVEL POLYNUCLECTIDES AND
FILL REFERENCE: 21402-114
CURRENT APPLICATION NUMBER: US/09/954,342
CURRENT FILING DATE: 2001-09-17
PRIOR APPLICATION NUMBER: 60/233,382
PRIOR FILING DATE: 2000-09-18
PRIOR FILING DATE: 2000-09-18
PRIOR FILING DATE: 2000-09-18
PRIOR PILING DATE: 2001-01-01
PRIOR APPLICATION NUMBER: 60/260,284
PRIOR APPLICATION NUMBER: 60/260,973
PRIOR APPLICATION NUMBER: 60/260,973
PRIOR APPLICATION NUMBER: 60/264,794
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/264,794
PRIOR FILING DATE: 2001-01-29
PRIOR APPLICATION NUMBER: 60/238,398
PRIOR APPLICATION NUMBER: 60/238,398
PRIOR APPLICATION NUMBER: 60/238,755
PRIOR APPLICATION NUMBER: 60/232,675
PRIOR APPLICATION NUMBER: 60/232,675
PRIOR FILING DATE: 2000-09-15
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US-09-954-342-65
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PRIOR FILING DATE: 2000-09-18
PRIOR PPLICATION NUMBER: 60/233,521
PRIOR FILING DATE: 2000-09-19
PRIOR APPLICATION NUMBER: 60/233,522
PRIOR APPLICATION NUMBER: 60/232,679
PRIOR APPLICATION NUMBER: 60/232,679
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APPLICANT: SPYTEK
APPLICANT: TAUPIE
APPLICANT: VERNET
APPLICANT: COLMAN
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Best Local Similarity
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APPLICANT:
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NUMBER OF SEQ ID NOS: 104
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PRIOR FILING DATE: 2000-09-15
PRIOR APPLICATION NUMBER: 60/233,960
PRIOR FILING DATE: 2000-09-20
PRIOR FILING DATE: 2000-09-20
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GERLACH, VALERIE L.
GERLACH, JOHN R.
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MACDOUGALL, UC...
MACDOUGALL, UC...
MACDOUGALL, UC...
MACDOUGALL, UC...
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SPYTEK, KIMBERLY ANN
TAUPIER, RAYMOND J.
                                                                                                                                                                                                                                                                                                                                                                                                                                         GUNTHER, ERIK
ELLERMAN, KAREN
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STONE, DAVID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              VERNET, CORINE A. COLMAN, STEVEN D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MILLET, ISABELLE
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FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 146
LENGTH: 104
TYPE: PRT
ORGANISM: Bos taurus
US-10-032-201B-146
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US-10-032-201B-146

US-10-032-201B-146, Application US/10032201B

Publication No. US20030167524A1
RESULT 27
US-10-032-201B-150
; Sequence 150, Application US/10032201B
; Publication No. US20030167524A1
; GENERAL INFORMATION:
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Best Local Similarity
Watches 6; Conserve
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Best Local Similarity
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PRIOR
PRIOR
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
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APPLICANT: Deckers, Harm
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PRIOR APPLICATION NUMBER: 60/232,679
PRIOR FILING DATE: 2000-09-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/233,801 FILING DATE: 2000-09-19
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Heifetz, Peter Bernard
Briggs, Steven
Dalmia, Bipin Kumar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Del Val, Greg
Zaplachinski, Steve
                                                                                                                                                                              100.0%; Score 47; DB ilarity 100.0%; Pred. No. 12; Conservative 0; Mismatches
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Pred. No. 12;
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APPLICANT: Van Rooijen, Gijs APPLICANT: Deckers, Harm APPLICANT: Heifetz, Peter Bernard

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RESULT 29
US-10-032-201B-154
; Sequence 154, Application US/10032201B
; Publication No. US20030167524A1
; GENERAL INFORMATION;
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US-10-032-201B-152
; Sequence 152, Application US/10032201B
; Publication No. US20030167524A1
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Best Local S
Matches 6
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 150
SEQ ID NO 150
LENGTH: 104
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SEQ ID NO 152
LENGTH: 104
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
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APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter B
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APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 38814 351B
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ORGANISM: Gallus gallus
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                                                                                                                                        30 WCGPCK 35
                                                                                                                                                                                                            ch 100.0%; Similarity 100.0%; 6; Conservative 0
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Heifetz, Peter Bernard
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Dalmia, Bipin Kumar
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Del Val, Greg
Zaplachinski, Steve
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                                                                                                                                                                                   RESULT 31
US-10-032-201B-158
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APPLICANT: Zaplachinski, Steve
APPLICANT: Zaplachinski, Steve
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 3518
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FREESEQ for Windows Version 4.0
SEQ ID NO 154
                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 155
SEQ ID NO 155
TENGTH: 104
TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                   Sequence 158, Application US/10032201B Publication No. US20030167524A1 GENERAL INFORMATION:
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Best Local (
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Best Local
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APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
                     APPLICANT:
                                                             APPLICANT:
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APPLICANT: Deckers, Harm
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APPLICANT: Deckers, Harm
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TYPE: PRT
ORGANISM: Macaca mulatta
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6; Conserva
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Heifetz, Peter Bernard
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Dalmia, Bipin Kumar
                                                    Deckers, Harm
Heifetz, Peter Bernard
                     Briggs, Steven
Dalmia, Bipin Kumar
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Greg
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 47; DB 4; Length 104; 100.0%; Pred. No. 12;
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                                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Rattus norvegicus US-10-032-201B-159
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 158
LENGTH: 104
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 313
SOPTWARE: FRANKSEQ for Windows Version 4.0
SEQ ID NO 159
LENGTH: 104
TYPE: PRT
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Matches
                                                                                                                                                                     Sequence 161, Appropriate Publication No.
                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                        GENERAL INFORMATION
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APPLICANT: Deckers, Harm
APPLICANT: Helfetz, Peter Bernard
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS,
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
                APPLICANT:
                                                                APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kun
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CURRENT FILING DATE: 2001-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
                                                    APPLICANT:
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les 6; Conserv
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Invention: Maurice
INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS,
                                                                                                                                                                                                                                                                                                              1 WCGPCK 6
                              Briggs, Steven
Dalmia, Bipin Kumar
Del Val, Greg
Zaplachinski, Steve
                                                                                                    Deckers, Harm
Heifetz, Peter Bernard
                                                                                                                                                                       Application US/10032201B
lo. US20030167524A1
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o. US20030167524A1
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Pred. No.
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     AND RELATED
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RESULT 35
US-10-424-599-282588
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US-10-032-201B-161
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Publication No. US20030167524A1
GENERAL INFORMATION:
APPLICANT: Van Rooijen, Gijs
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Matches
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 161
LENGTH: 104
APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 202588
LENGTH: 104
                                                                                                                                                                                                                                  Sequence 282588, Application US/10424599 Publication No. US20040031072A1 GENERAL INFORMATION:
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SEQ ID NO 276
LENGTH: 104
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 3518
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Staphylococcus aureus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
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Pred. No.
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Pred. No.
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PRIOR APPLICATION NUMBER: EP 01870003.9
PRIOR FILING DATE: 2001-01-09
NUMBER OF SEQ ID NOS: 732
SOFTWARE: PatentIn version 3.1
SEQ ID NO 188
LENGTH: 104
TYPE: PRT
ORGANISM: Saccharomyces cerevisiae
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US-09-954-342-61
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APPLICANT: MISHRA, VISHNU S.
APPLICANT: SPYTEK, KIMBERLY
APPLICANT: TAUPIER, RAYMOND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Publication No.
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Best Local Similarity
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APPLICANT:
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TITLE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION IN
TITLE OF INVENTION: YEAST AND FUNGI
FILE REFERENCE: JAB-1667
CURRENT APPLICATION NUMBER: US/10/451,467A
CURRENT FILING DATE: 2003-06-19
PRIOR APPLICATION NUMBER: EP 00870318.3
PRIOR FILING DATE: 2001-12-22
PRIOR APPLICATION NUMBER: EP 01870002.1
PRIOR FILING DATE: 2001-01-04
PRIOR FILING DATE: 2001-01-04
PRIOR FILING DATE: 2001-01-04
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APPLICANT:
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                                                                                                                                                                                                                                                                                   Application US/09954342
                                                               PADIGARU, MURALIDHARA
GERLACH, VALERIE L.
MACDOUGALL, JOHN R.
                                                                                                                                                               VERNET, CORINE A.
COLMAN, STEVEN D.
GORMAN, LINDA
                                                                                                                                                                                                                 SPYTEK, KIMBERLY ANN
TAUPIER, RAYMOND J.
 STONE,
                                               MACDOUGALL, JOHN I
SMITHSON, GLENNDA
                                                                                                                  SHENOY, SURESH
                                                                                                                                TCHERNEV, VELIZAR T.
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PEYMAN, JOHN
STONE, DAVID
                             MILLET, ISABELLE
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o. US20040161840A1
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RESULT 38
US-10-171-311-226
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Publication No. US20030087270A1
GENERAL INFORMATION:
APPLICANT: Schlegel, Robert
APPLICANT: Chen, Yan
APPLICANT: Zhao, Xumei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver.
SEQ ID NO 61
LENGTH: 105
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Best Local :
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               APPLICANT: Monahan, John
APPLICANT: Kamatkar, Shubhangi
APPLICANT: Glatt, Karen
APPLICANT: Glatt, Karen
APPLICANT: Gennavarapu, Manjula
APPLICANT: Hoersh, Sebastian
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS, KITS, AND METHODS
TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND
TITLE OF INVENTION: OF CERVICAL CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/260,973
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/264,794
PRIOR FILING DATE: 2001-01-29
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PRIOR APPLICATION NUMBER: 60/233,382
PRIOR FILING DATE: 2000-09-18
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NUMBER OF SEQ ID NOS: 104
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FILE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
nes 6; Conserv
REFERENCE: MRI-035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/233,801
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APPLICATION NUMBER: 60/274,862
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Pred. No.
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Mismatches
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                                                                    GENERAL INFORMATION:
APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Ber
APPLICANT: Briggs, Steven
APPLICANT: Briggs, Steven
APPLICANT: Del Val, Greg
APPLICANT: Del Val, Greg
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SEQ ID NO 226
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LENGTH: 105
                                                                                                                                                                                                       Sequence 47, Appropriate Publication No.
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CURRENT FILING DATE: 2002-12-10
PRIOR APPLICATION NUMBER: US 60/355,295
PRIOR FILING DATE: 2002-02-08
NUMBER OF SEQ ID NOS: 308
SOFTWARE: PatentIn version 3.1
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PRIOR APPLICATION NUMBER: US 60/298,159
PRIOR FILING DATE: 2001-06-13
PRIOR APPLICATION NUMBER: US 60/298,155
PRIOR FILING DATE: 2001-06-13
PRIOR APPLICATION NUMBER: US 60/335,936
PRIOR APPLICATION NUMBER: US 60/335,936
PRIOR FILING DATE: 2001-11-14
                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Greis, Kenneth
TITLE OF INVENTION: Angiogenesis Modulating Proteins
FILE REFERENCE: 8865M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: The Procter & Gamble Company
APPLICANT: Peters, Kevin
APPLICANT: Thompson, Larry
APPLICANT: Wang, Feng
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SOFTWARE: FastSEQ for Windows Version 4.0
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ORGANISM: Rattus norvegicus
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TYPE: PRT
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les 6; Conserv
OF INVENTION:
                                                                                                                                                                                                                                                                                                                      31 WCGPCK 36
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I: Moloney, Maurice INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                         1 WCGPCK 6
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                                                                                     Deckers, Harm
Heifetz, Peter Bernard
Briggs, Steven
Dalmia, Bipin Kumar
                                                                                                                                                                                                       Application US/10032201B
No. US20030167524A1
                                                      Del Val, Greg
Zaplachinski, Steve
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100.0%; Pred. No.
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Pred. No.
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FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 106
LENGTH: 105
TYPE: PRT
ORGANISM: Cornybacterium nephridii
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US-10-032-201B-106
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APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Chou Yihua
APPLICANT: Chou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 237784
LENGTH: 105
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 47
LENGTH: 105
TYPE: PRT
ORGANISM: Homo Sapien
                                                                                                                                                                                                                                       Sequence 237784, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
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TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
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Dalmia, Bipin Kumar
Del Val, Greg
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100.0%; Pred. No.
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SOFTWARE: PASTSEQ for Windows Version 4.0
SEQ ID NO 127
LENGTH: 105
TYPE: PRT
ORGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank BAA04881
DATABASE ENTRY DATE: 2002-12-25
US-10-351-891-127
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                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Ghosh, Sc
APPLICANT: Fahy, Ec
APPLICANT: Zhang, E
SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2019 LENGTH: 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 127, Application US/10351891 Publication No. US20040048311A1
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Matches
                                                     APPLICANT: Gibson, Bradford W.
APPLICANT: Taylor, Steven W.
APPLICANT: Glenn, Gary M.
APPLICANT: Glenn, Gary M.
APPLICANT: Warnock, Dale E.
TITLE OF INVENTION: TARGETS FOR THERAPEUTIC INTERVENTION
TITLE OF INVENTION: IDENTIFIED IN THE MITOCHONDRIAL PROTEOME
FILE REFERENCE: 660088.465
CURRENT APPLICATION NUMBER: US/10/408,765A
CURRENT FILING DATE: 2003-04-04
NUMBER OF SEQ ID NOS: 3077
CONTRADES FOREORE FOREORE
                                                                                                                                                                                                                                                                                                                                                                  Sequence 2019, Application US/10408765A Publication No. US20040101874A1
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TITLE OF INVENTION: USE OF COLLECTIONS OF BINDING
TITLE OF INVENTION: APPLICATIONS
FILE REFERENCE: 25885-1753
CURRENT APPLICATION NUMBER: US/10/351,891
CURRENT FILING DATE: 2003-01-24
PRIOR APPLICATION NUMBER: US 60/352,011
PRIOR PILING DATE: 2002-01-24
NUMBER OF SEO ID NOS: 140
NUMBER OF SEO ID NOS: 140
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LOCATION: (1)..(105)
OTHER INFORMATION: u
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6; Conserv
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                                                                                                                                                                                                                                                                               Zhang, Bing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Score 47; DB ilarity 100.0%; Pred. No. 12; Conservative 0; Mismatches
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ilarity 100.0%; Pred. No. 12
Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Length 105;
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Query Match
Best Local Similarity
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 RESULT 45
US-10-660-118A-8
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 ; ORGANISM: Homo sapiens
US-10-408-765A-2019
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 US-10-660-118A-9
 US-10-660-118A-9
 RESULT 46
 US-10-660-118A-8
 Sequence 9, Application US/10660118A
Publication No. US20040131606A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum
FILE REFERENCE: 2879-98
 SOFTWARE: PatentIn version 3.1
SEQ ID NO 8
LENGTH: 105
 Sequence 8, Application US/10660118A
Publication No. US20040131606A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
 NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.1
SEQ ID NO 9
 Matches
 Query Match
Best Local :
 Matches
 Query Match
Best Local Similarity
 CURRENT APPLICATION NUMBER: US/10/660,118A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
 FILE REFERENCE: 2879-98
CURRENT APPLICATION NUMBER: US/10/660,118A
CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
 PRIOR FILING DATE: 2003-
NUMBER OF SEQ ID NOS: 15
 TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum
 PRIOR FILING DATE:
 TYPE: PRT
ORGANISM: Mus musculus
 TYPE: PRT
ORGANISM: Gallus gallus
 LENGTH: 105
 31 WCGPCK 36
 31 WCGPCK 36
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 1 WCGPCK 6
 1 WCGPCK 6
 1 WCGPCK 6
 Similarity 6; Conserv
 6,
WCGPCK 36
 Conservative
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 2003-04-11
 2003-04-11
 100.0%; Score 47;
100.0%; Pred. No.
 100.0%;
 100.0%; Score 47; DB 4; 100.0%; Pred. No. 12;
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 Score 47;
Pred. No.
 Mismatches
 Mismatches
 DB 4;
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 Length 105
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 RESULT 49
US-10-660-118A-12
 US-10-660-118A-10
 US-10-660-118A-10
 RESULT 47
 US-10-660-118A-11
 US-10-660-118A-11
 RESULT 48
Sequence 12, Application US/10660118A
Publication No. US20040131606A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum
FILE REFERENCE: 2879-98
CURRENT APPLICATION NUMBER: US/10/660,118A
 Sequence 11, Application US/10660118A Publication No. US20040131606A1 GENERAL INFORMATION:
 Sequence 10, Application US/10660118A
Publication No. US20040131606A1
GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.1 SEQ ID NO 11
 SEQ ID NO 10
 APPLICANT: White, Carl W.

APPLICANT: White, Carl W.

TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum FILE REFERENCE: 2879-98

CURRENT APPLICATION NUMBER: US/10/660,118A

CURRENT FILING DATE: 2003-09-10

PRIOR APPLICATION NUMBER: 60/409,960

PRIOR APPLICATION NUMBER: 60/462,082

PRIOR APPLICATION NUMBER: 60/462,082
 Query Match
Best Local Similarity
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 Matches
 Query Match
 CURRENT APPLICATION NUMBER: US/10/660,118A CURRENT FILING DATE: 2003-09-10 PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10 PRIOR APPLICATION NUMBER: 60/462,082
 PRIOR FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.1
 APPLICANT: White, Carl W.
TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum
FILE REFERENCE: 2879-98
 PRIOR FILING DATE: 2003-
NUMBER OF SEQ ID NOS: 15
 TYPE: PRT
ORGANISM: Rattus norvegicus
 ORGANISM: Bos taurus
 TYPE: PRT
 LENGTH: 105
 LENGTH:
 Local Similarity
 31 WCGPCK 36
 31 WCGPCK 36
 6,
 105
 1 WCGPCK 6
 1 WCGPCK 6
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 Conservative
 Carl W.
 2003-04-11
 100.0%;
 100.0%; Score 47; DB
100.0%; Pred. No. 12;
tive 0; Mismatches
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 Score 47;
Pred. No.
 Mismatches
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CURRENT FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/409,960

PRIOR FILING DATE: 2002-09-10

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RESULT 51
US-10-699-113-32
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 RESULT 50
US-10-451-467A-684
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 ; TYPE: PRT; ORGANISM: Homo sapiens US-10-660-118A-12
 US-10-451-467A-684
 SOFTWARE: PatentIn version 3.1
SEQ ID NO 12
LENGTH: 105
 Sequence 32, Application US/10699113 Publication No. US20040241748A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 SEQ ID NO 684
LENGTH: 105
 Sequence 684, Application US/10451467A Publication No. US20040161840A1 GENERAL INFORMATION:
 Matches
 Matches
 Query Match
Best Local Similarity
 APPLICANT: Ault-Riche, Dana
APPLICANT: Kumble, Krishnanand
APPLICANT: Schulz, Rainer
APPLICANT: Schulz, Kenneth
TITLE OF INVENTION: Self-Assembling Arrays and Uses Thereof
FILE REFERENCE: 25885-1755
 CURRENT APPLICATION NUMBER: US/10/451,467A
CURRENT FILING DATE: 2003-06-19
PRIOR APPLICATION NUMBER: EP 00870318.3
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: EP 01870002.1
PRIOR FILING DATE: 2001-01-04
 CURRENT APPLICATION NUMBER: US/10/699,113
CURRENT FILING DATE: 2003-10-30
 APPLICANT: REEKWANS, RIEKA JOSEPHINA
TITLE OF INVENTION: BAX-RESPONSIVE GENES FOR DRUG TARGET IDENTIFICATION IN
TITLE OF INVENTION: YEAST AND FUNGI
FILE REFERENCE: JAB-1667
 PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
NUMBER OF SEQ ID NOS: 15
 NUMBER OF SEQ ID NOS: 732
 SOFTWARE: PatentIn version 3.1
 PRIOR FILING DATE: 2001-01-09
 PRIOR APPLICATION NUMBER: EP 01870003.9
 APPLICANT:
 APPLICANT: CONTRERAS, ROLAND HENRI
PRIOR APPLICATION NUMBER: 60/446,687
 TYPE: PRT
ORGANISM: Homo sapiens
 31 WCGPCK 36
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 In 100.0%; Score 47; DB 4; Length 105; Similarity 100.0%; Pred. No. 12; 6; Conservative 0; Mismatches 0.
 6;
 EBERHARDT, INES
LUYTEN, WALTER HERMAN MARIA LOUIS
 Conservative
 100.0%; Score 47; DB 4; 100.0%; Pred. No. 12;
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 Mismatches
 Length 105;
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 Gaps
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; TYPE: PRT ; ORGANISM: Macaca mulatta US-10-970-370-6
 RESULT 53
US-10-970-370-6
 US-10-970-370-2
 US-10-970-370-2
 US-10-699-113-32
 NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 105
 Sequence 2, Application US/10970370 Publication No. US20050208037A1 GENERAL INFORMATION:
APPLICANT: DASHNAMOORTHY, RAVI APPLICANT: DAS, KUMUDA C.
 PRIOR FILING DATE: 2003-02-10
NUMBER OF SEQ ID NOS: 948
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ:ID:NO 32
 SOFTWARE: PatentIn Ver. SEQ ID NO 6
 GENERAL INFORMATION:
APPLICANT: DASHNAMOORTHY, RAVI
APPLICANT: DAS, KUMUDA C.
 Matches
 Sequence 6, Application US/10970370 Publication No. US20050208037A1
 Query Match 100.0%; Score 47; DB Best Local Similarity 100.0%; Pred. No. 12;
 Matches
 Query Match
Best Local Similarity
 TITLE OF INVENTION: THIOREDOXIN INCREASES REDOX-CYCLING OF A TITLE OF INVENTION: AGENTS THEREBY SENSITIZES CANCER CELLS FILE REFERENCE: UTSN:025US CURRENT APPLICATION NUMBER: US/10/970,370 CURRENT FILING DATE: 2004-10-21 PRIOR APPLICATION NUMBER: 60/513,134 PRIOR APPLICATION NUMBER: 60/513,134 PRIOR APPLICATION DATE: 2003-10-21 NUMBER: 00/510,003: 9
 CURRENT APPLICATION NUMBER: US/10/970,370
CURRENT FILLING DATE: 2004-10-21
PRIOR APPLICATION NUMBER: 60/513,134
PRIOR FILING DATE: 2003-10-21
 TITLE OF INVENTION: THIOREDOXIN INCREASES REDOX-CYCLING OF ANTICANCER TITLE OF INVENTION: AGENTS THEREBY SENSITIZES CANCER CELLS TO APOPTOSIS FILE REFERENCE: UTSN:025US
 PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank BAA04881
DATABASE ENTRY DATE: 2002-12-25
 LENGTH: 105
TYPE: PRT
ORGANISM: Homo sapiens
 TYPE: PRT
ORGANISM: Mus musculus
 LENGTH: 105
 ENGTH: 105
 31 WCGPCK 36
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 100.0%; l Similarity 100.0%; 6; Conservative 0
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 Score 47;
Pred. No.
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 DB 5; Length 105; 12;
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 Query Match
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 US-10-032-201B-132
 US-10-032-201B-123
 RESULT 55
 US-10-032-201B-123
 APPLICANT: MOIONEY, MAURICE
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 123
LENGTH: 106
 NUMBER OF SEQ ID NOS: 313
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 132
LENGTH: 106
 Sequence 132, Application US/10032201B Publication No. US20030167524A1
 Query Match
Best Local Similarity
 Query Match
 GENERAL INFORMATION
 APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: MCTHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 3518
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
 APPLICANT:
 Publication No. US20030167524A1
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT:
 TYPE: PRT ORGANISM: Penicillium chrysogenum
 TYPE: PRT
ORGANISM: Helicobacter pylori
 y Match 100.0%; Score 47; DB Local Similarity 100.0%; Pred. No. 12; hes 6; Conservative 0; Mismatches
 INFORMATION:
 31 WCGPCK 36
 29 WCGPCK 34
 1 WCGPCK 6
 1 WCGPCK 6
 Deckers, Harm
Heifetz, Peter Bernard
Briggs, Steven
Dalmia, Bipin Kumar
Del Val, Greg
Zaplachinski, Steve
 Deckers, Harm
Heifetz, Peter Bernard
 Application US/10032201B
 Conservative
 100.0%;
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 Score 47;
Pred. No.
 Score 47; DB Pred. No. 12;
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 176
LENGTH: 106
TYPE: PRT
ORGANISM: Manduca Bexta
US-10-032-201B-176
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 US-10-032-201B-176
 RESULT 57
 ; ORGANISM: Streptomyces aureofaciens US-10-032-201B-137
 US-10-032-201B-137
 RESULT 56
 Query Match
Best Local S
 CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
RUMBER OF SEQ ID NOS: 313
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 137
LENGTH: 106
 Sequence 137, Application US/10032201B Publication No. US20030167524A1
 Matches
 GENERAL INFORMATION:
 Sequence 176, Application US/10032201B Publication No. US20030167524A1
 Matches
 Query Match
Best Local &
 GENERAL INFORMATION
 Matches
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
APPLICANT: Briggs, Steven
 APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bo
 APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT:
 APPLICANT:
 APPLICANT:
 FILE REFERENCE: 38814 351B
 TYPE: PRT
 30 WCGPCK 35
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 Similarity 6; Conserva
 Similarity 6; Conserv
 6; Conservative
 Deckers, Harm
Heifetz, Peter Bernard
 Briggs, Steven
Dalmia, Bipin Kumar
 Deckers, Harm
Heifetz, Peter Bernard
 Briggs, Steven
Dalmia, Bipin Kumar
 Del Val, Greg
Zaplachinski, Steve
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 100.0%; Score 47;
100.0%; Pred. No.
 100.0%; Score 47; DB 100.0%; Pred. No. 12;
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RESULT 59
US-10-628-391-1
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 US-10-335-977-6752
 US-10-335-977-6752
 Sequence 1, Application US/10628391
Publication No. US20040146528A1
GENERAL INFORMATION:
APPLICANT: WINDLE, Henry J.
APPLICANT: O'TOOLE, Dermot
APPLICANT: KELLEHOR, Dermot
APPLICANT: KELLEHOR, Dermot
APPLICANT: KELLEHOR, Dermot
APPLICANT: KELLEHOR.
 Sequence 6752, Application US/10335977

Publication No. US20040052799A1

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: DOUGLAS SMITH et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES
RELATING TO HELICOBACTER PYLORI FOR
DIAGNOSTICS AND THERAPEUTICS
 Matches
 Query Match
TITLE OF INVENTION: INHIBITION OF NF-kappaB ACTIVATION
 TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO: 6752:
SEQUENCE CHARACTERISTICS:
 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/335,977
FILING DATE: 30-Dec-2002
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/993,002
FILING DATE: 17-DEC-1997
ATTORNEY/AGENT INFORMATION:
NAME: Mandragouras, Amy E.
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: GTN-018
TELECOMMUNICATION INFORMATION:
 Local Similarity 100 nes 6; Conservative
 SOPTWARE: UNIX
 NAME/KEY: misc_feature
LOCATION: (B) LOCATION 1...106
SEQUENCE DESCRIPTION: SEQ ID NO: 6752:
 HYPOTHETICAL: YES
ORIGINAL SOURCE:
ORGANISM: Helicobacter pylori
 COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: Windows NT 4.0
 FEATURE:
 MOLECULE TYPE: protein
 CORRESPONDENCE ADDRESS:
 NUMBER OF SEQUENCES: 10031
 29
 1 WCGPCK 6
 WCGPCK 34
 LENGTH: 106 amino acids TYPE: amino acid TOPOLOGY: linear
 COUNTRY: USA
 STATE: Massachusetts
 CITY: Boston
 STREET: 28 State Street
 TELEPHONE: (617)227-7400
 ADDRESSEE: LAHIVE & COCKFIELD
 02109-1875
 100.0%;
 0; Mismatches
 Score 47;
Pred. No.
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 0
 Length 106;
 Indels
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 Gaps
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APPLICANT: Venter, J. Craig

APPLICANT: et al.

TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID

TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE

TITLE OF INVENTION: DROSOPHILA GENES.

FILE REFERENCE: CL000728

CURRENT APPLICATION NUMBER: US/11/097,143

CURRENT FILING DATE: 2005-04-04

PRIOR APPLICATION NUMBER: 60/157,832

PRIOR APPLICATION NUMBER: 60/150,191

PRIOR APPLICATION NUMBER: 60/160,191

PRIOR APPLICATION NUMBER: 60/161,932

PRIOR APPLICATION NUMBER: 60/161,932

PRIOR FILING DATE: 1999-10-19

PRIOR FILING DATE: 1999-10-28
 US-11-097-143-7479
Sequence 7479, Application US/11097143
Publication No. US20050208558A1
GENERAL INFORMATION:
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 ; OTHER INFORMATION: Clone ID: MRT4577_96874C.1.pep US-10-425-115-365910
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 Query Match
Best Local Similarity
Thes 6; Conserve
 ; TYPE: PRT; ORGANISM: H. pylori
US-10-628-391-1
 RESULT 61
 US-10-425-115-365910
 Query Match
Best Local S
Matches 6
 NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 365910
LENGTH: 106
 GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
 SOFTWARE: PatentIn version 3.2 SEQ ID NO 1
 Sequence 365910, Application US/10425115 Publication No. US20040214272A1
 TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants FILE REFERENCE: 38-21(53222)B CURRENT APPLICATION NUMBER: US/10/425,115 CURRENT FILING DATE: 2003-04-28
 CURRENT APPLICATION NUMBER: US/10/628,391
CURRENT FILING DATE: 2003-07-29
PRIOR APPLICATION NUMBER: PCT/IE02/00011
PRIOR FILING DATE: 2002-01-30
NUMBER OF SEQ ID NOS: 9
 FILE REFERENCE: P69048US0
 TYPE: PRT
ORGANISM: Zea mays
FEATURE:
 ENGTH:
 Local Similarity
 29 WCGPCK 34
 31 WCGPCK 36
 106
 1 WCGPCK 6
 1 WCGPCK 6
 Conservative
 Conservative
 100.0%;
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 Score 47; DB
Pred. No. 12;
 Score 47; DB Pred. No. 12;
 Mismatches
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 DB 4; Length 106;
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 Length 106
 Indels
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PRIOR APPLICATION NUMBER: 60/173,383
PRIOR FILING DATE: 1999-12-28
PRIOR APPLICATION NUMBER: 60/175,693
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/184,831
PRIOR PILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: 60/191,637
PRIOR APPLICATION NUMBER: 60/191,637
PRIOR PILING DATE: 2000-03-23
NUMBER OF SEQ ID NOS: 43008
SOFTWARE: FBSSECSEQ for Windows Version 4.
SEQ ID NO 7479
LENGTH: 106
 RESULT 63
US-10-032-201B-121
; Sequence 121, Application US/10032201B
; Publication No. US20030167524A1
; GENERAL INFORMATION:
; APPLICANT: Van ROOijen, Gijs
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 US-10-032-201B-109
 US-10-032-201B-109
 ; ORGANISM: DROSOPHILA US-11-097-143-7479
 CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 109
LENGTH: 107
 Matches
 GENERAL INFORMATION:
 Publication No.
 Query Match
Best Local S
 Query Match
Best Local :
 APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter B
APPLICANT:
 FEATURE: VARIANT
NAME/KEY: VARIANT
LOCATION: 17, 38, 42, 55, 58, 60, 72, 107
OTHER INFORMATION: Xaa = Any Amino Acid
 ORGANISM: Chromatium vinosum
 TYPE: PRT
 y match 100.0%;
Local Similarity 100.0%;
hes 6; Conservative 0,
 Local Similarity nes 6; Conserv
 FILING DATE: 1999-11-
 APPLICATION NUMBER: 60/164,769
 31 WCGPCK 36
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 Deckers, Harm
Heifetz, Peter Bernard
 Application US/10032201B o. US20030167524A1
 Conservative
 100.0%; Score 47; DB
100.0%; Pred. No. 12;
Live 0; Mismatches
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 Score 47; DB 6; Length 106; Pred. No. 12;
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 RESULT 64
 Matches
 Query Match
Best Local :
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 Query Match
Best Local (
 TYPE: PRT
 LENGTH: 107
 LENGTH: 107
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; TYPE: PRT
; ORGANISM: Haemophilus influenzae
US-10-032-201B-121
 RESULT 65
US-10-335-977-6753
 ; ORGANISM: Drosophila melanogaster US-10-032-201B-151
 US-10-032-201B-151
 Sequence 6753, Application US/10335977
Publication No. US20040052799A1
GENERAL INFORMATION:
APPLICANT: DOUGLAS SMITH et al
TITLE OF INVENTION: NUCLEIC ACII
 FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 121
 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 151
 Sequence 151, A Publication No.
 GENERAL INFORMATION:
 APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Bernard
APPLICANT: Briggs, Steven
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 FILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
 APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
NUMBER OF SEQUENCES:
 30 WCGPCK 35
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 Similarity 6; Conserv
 Similarity 6; Conserv
 Heifetz, Peter Bern
Briggs, Steven
Dalmia, Bipin Kumar
 Zaplachinski,
 Application US/10032201B
o. US20030167524A1
 Conservative
 Conservative
 DOUGLAS SMITH et al
NVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES
NVENTION: RELATING TO HELICOBACTER PYLORI FOR
DIAGNOSTICS AND THERAPEUTICS
 Peter Bernard
 100.0%;
 100.0%;
 Steve
 10031
 0;
 0,
 Score 47; DB
Pred. No. 12;
0; Mismatches
 Score 47;
Pred. No.
 Mismatches
 DB
12;
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 4; Length 107;
 4; Length 107;
 Indels
 Indels
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 <u>,</u>
 Gaps
 Gaps
 <u>,,</u>
 0
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CORRESPONDENCE ADDRESS:

COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660

02109-1875

STATE: Massachusetts CITY: Boston

STREET: 28 State Street

ADDRESSEE: LAHIVE & COCKFIELD

SOFTWARE: UNIX

COMPUTER: IBM PC Compatible OPERATING SYSTEM: Windows NT 4.0

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Sequence 8226, Application US/11097143

| Publication No. US20050208558A1 |
| GENERAL INFORMATION: | GENERAL INFORMATION: | APPLICANT: Venter, J. Craig |
| APPLICANT: Venter, J. Craig |
| APPLICANT: et al. |
| TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID |
| TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID |
| TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID |
| TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID |
| TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID |
| TITLE OF INVENTION: DETECTION ACROSCOPHILA GENES. |
| TITLE OF INVENTION: DETECTION OF 10,000 OR MORE |
| TITLE OF INVENTION: DETECTION OF 10,000 OR MORE |
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LOCATION: (B) LOCATION 1...107;
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SEQUENCE DESCRIPTION: SEQ ID NO: 6753;
US-10-335-977-6753
 RESULT 66
US-11-097-143-8226
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Best Local Similarity lou.
 TELEPHONE: (617)227-7400
TELEPAX: (617)742-4214
INFORMATION FOR SEQ ID NO: 6753:
SEQUENCE CHARACTERISTICS:
 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/335,977

FILING DATE: 30-Dec-2002

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/993,002

FILING DATE: 17-DEC-1997

ATTORNEY/AGENT INFORMATION:

NAME: Mandragouras, Amy E.

REGISTRATION NUMBER: 36,207

REFERENCE/DOCKET NUMBER: GTN-018

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
FILING DATE: 1999-11-12
APPLICATION NUMBER: 60/173,383
FILING DATE: 1999-12-28
 HYPOTHETICAL: YES ORIGINAL SOURCE:
 FEATURE:
 MOLECULE TYPE: protein
 30 WCGPCK 35
 1 WCGPCK 6
 LENGTH: 107 amino acids
TYPE: amino acid
TOPOLOGY: linear
 ORGANISM: Helicobacter pylori
 100.0%;
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 Score 47; DB
Pred. No. 12;
 4; Length 107;
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PRIOR FILING DATE: 2000-01-12
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PRIOR FILING DATE: 2000-03-23
NUMBER OF SEQ ID NOS: 43008
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8226
LENGTH: 107
TYPE: PRT
 Sequence 117, Application US/10
Publication No. US/20030167524A1
GENERAL INFORMATION:
APPLICANT: Van Rooijen, Gijs
APPLICANT: Ueckers, Harm
APPLICANT: Briggs, Steven
APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kuman
 RESULT 68
US-10-032-201B-117
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 ; TYPE: PRT; ORGANISM: Escherichia coli US-09-812-777-16
 RESULT 67
US-09-812-777-16
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 ; ORGANISM: DROSOPHILA US-11-097-143-8226
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Matches 6
 CURRENT APPLICATION NUMBER: US/09/812,777
CURRENT FILING DATE: 2001-03-18
PRIOR APPLICATION NUMBER: KR 00-63596
PRIOR FILING DATE: 2000-10-27
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 16
PILE REFERENCE: 38814 351B
CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 117
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 Sequence 16,
 Sequence 16, Application US/09812777 Patent No. US20020115827A1
 APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT: KIM, KEY-SUN
APPLICANT: LEB, DUCK YEON
TITLE OF INVENTION: A METHOD OF PROTEIN STABILIZATION
PILE REFERENCE: 05823.0195-00000
 ENGTH: 108
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 30 WCGPCK 35
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 1 WCGPCK 6
 1 WCGPCK 6
 Briggs, Steven
Dalmia, Bipin Kumar
 Heifetz, Peter Bernard
 Del Val, Greg
Zaplachinski, Steve
 Application US/100322018
 US20030167524A1
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 Gaps
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APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Moloney, Maurice
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
FILE REFERENCE: 38814 351B
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CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 13
LENGTH: 108
TYPE: PRT
ORGANISM: Pseudomonas aeruginosa
US-10-032-201B-133
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 RESULT 69
US-10-032-2018-133
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 ; TYPE: PRT ; ORGANISM: E coli, salmonella typhimurium US-10-032-201B-117
 US-10-032-201B-142
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CURRENT APPLICATION NUMBER: US/10/032,201B
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 313
SOFTWARE: FastSEQ for Windows Version 4.0
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 Sequence 133, Application US/10032201B Publication No. US20030167524A1
 APPLICANT: Briggs, Steven
APPLICANT: Dalmia, Bipin Kumar
APPLICANT: Del Val, Greg
APPLICANT: Zaplachinski, Steve
APPLICANT: Moloney, Maurice
APPLICANT: Moloney, Maurice
TITLE OF INVENTION: METHODS FOR THE PRODUCTION OF MULTIMERIC PROTEINS, AND RELATED
TITLE OF INVENTION: COMPOSITIONS
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Petter Be
APPLICANT: Briggs, Steven
 APPLICANT: Van Rooijen, Gijs
APPLICANT: Deckers, Harm
APPLICANT: Heifetz, Peter Be
 LENGTH: 108
ORGANISM: Thiobacillus ferrooxidans
 INFORMATION:
 31 WCGPCK 36
 32 WCGPCK 37
 1 WCGPCK 6
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Heifetz, Peter Bernard
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Heifetz, Peter Bernard
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US-10-032-201B-142

Query Match

Query Match

Best Local Similarity 100.0%; Pred. No. 12;

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Db 32 WCGPCK 37
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Search completed: Pebruary 23, 2006, 00:47:55 Job time : 125.5 Becs

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 Minimum DB
Maximum DB
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Perfect score:
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MASS

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Sequence 3, Application US/11090916

Publication No. US20050260140A1

GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: Well, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1

CURRENT APPLICATION NUMBER: US/11/090,916

CURRENT APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR PRIOR APPLICATION NUMBER: 60/462,082
PRIOR RILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR FILING DATE: 2005-02-07
NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
NUMBER: FOR SEQ ID NOS: 15
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 US-11-090-916-3
 SOFTWARE: PatentIn version 3.3
SEQ ID NO 3
LENGTH: 6
 ORGANISM: Artificial sequence FEATURE:
 OTHER INFORMATION: synthetic peptide motif
 TYPE: PRT
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 Sequence 2. Application US/11131744

Publication No. US20060023808A1

GENERAL INFORMATION:
APPLICANT: HAJIVANDI, MAHBOD R.
APPLICANT: SADEGHI, MEHRNOOSH
APPLICANT: LIANG, XIQUAN
COURTENT APPLICATION COMPOSITIONS, KITS, AND METHODS FOR CALIBRATION IN MASS
TILLE OF INVENTION: SPECTROMETRY
FILE REFERENCE: NUX-28 US
CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT APPLICATION NUMBER: 60/572,290
PRIOR APPLICATION NUMBER: 60/572,290
PRIOR APPLICATION NUMBER: 60/59,373
PRIOR APPLICATION NUMBER: 60/69,373
PRIOR APPLICATION NUMBER: 60/69,373
PRIOR APPLICATION NUMBER: 60/69,373
PRIOR FILING DATE: 2005-04-08
PRIOR APPLICATION NUMBER: 60/691,685
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR APPLICATION NUMBER: 60/621,686
TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR CALIBRATION IN MASS
TITLE OF INVENTION: SPECTROMETRY
FILE REFERENCE: NVX.-28 US
CURRENT APPLICATION UNMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR APPLICATION NUMBER: 60/672,215
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR APPLICATION NUMBER: 60/621,685
PRIOR FILING DATE: 2004-05-17
PRIOR APPLICATION NUMBER: 60/621,686
PRIOR APPLICATION NUMBER: 60/621,686
PRIOR APPLICATION NUMBER: 60/621,686
PRIOR APPLICATION NUMBER: 60/621,686
PRIOR PILING DATE: 2004-10-26
NUMBER: 00/621,686
PRIOR PILING DATE: 2004-10-26
SOPTWARE: Patentin Ver. 3.3
SEQ ID NO 1
LENGTH: 91
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 US-11-131-744-1
 US-11-131-744-2
 US-11-131-744-2
 Sequence 1, Application US/11131744
Publication No. US20060023808A1
GENERAL INFORMATION:
 SOFTWARE: PatentIn Ver. 3.3
SEQ ID NO 2
LENGTH: 75
TYPE: PRT
ORGANISM: Artificial Sequence
 Matches
 Query Match
Best Local Similarity
 APPLICANT: HAJIVANDI, MAHBOD R.
APPLICANT: POPE, ROBERT M.
APPLICANT: SADEGHI, MEHRNOOSH
APPLICANT: LIANG, XIQUAN
APPLICANT: LEITE, JOHN F.
 OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
 FEATURE:
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 Gaps
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; TYPE: PRT; ORGANISM: Artificial Sequence; FEATURE: INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: protein sequence
US-11-131-744-1
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 ; TYPE: PRT
; ORGANISM: Lactobacillus acidophilus
US-11-074-176-286
 US-11-090-916-7
 US-11-074-176-286
 Sequence 7, Application US/11090916 Publication No. US20050260140A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 SEQ ID NO 286
LENGTH: 103
 Sequence 286, Application US/11074176 Publication No. US20050250135A1
 Matches
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Best Local Similarity
 APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
 GENERAL INFORMATION:
 PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
 PRIOR APPLICATION NUMBER: 60/551,161
PRIOR FILING DATE: 2004-03-08
NUMBER OF SEQ ID NOS: 381
 FILE REFERENCE: 5051-694
CURRENT APPLICATION NUMBER: US/11/074,176
CURRENT FILING DATE: 2005-03-07
 APPLICANT: Peril, Andrea Azcarate
TITLE OF INVENTION: Nucleic Acid Sequences Encoding
TITLE OF INVENTION: Stress-Related Proteins and Uses Therefore
 APPLICANT: Klaenhammer, Todd R. APPLICANT: Russell, William M.
 APPLICANT:
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 SOFTWARE: FastSEQ for Windows Version 4.0
APPLICATION NUMBER: 60/650,865
 FILING DATE: 2004-03-24
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McAuliffe, Olivia
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APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
ITILE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2005-02-07
NUMBER: OF SEQ ID NOS: 15
PRIOR FILING DATE: 2005-02-07
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US-10-821-234-1371
 RESULT 6
 US-10-821-234-1371
 NUMBER OF SEQ ID NOS: 1704
SOFTWARE: pt_SEQ_genes Version 1.0
SEQ ID NO 1371
FEBRUARY
 Sequence 8, Application US/11090916 Publication No. US20050260140A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 7
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 Sequence 1371, Publication No.
 GENERAL INFORMATION:
 FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILLING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
 APPLICANT: Labat, Ivan APPLICANT: Stache-Cra
 APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
 TYPE: PRT
 ENGTH: 105
 Local Similarity
 1371, Application US/10821234 ion No. US20050255114A1
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 US-11-090-916-9
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
 Sequence 10, Application US/11090916 Publication No. US20050260140A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.3
SEQ ID NO 9
LENGTH: 105
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Best Local Similarity
 GENERAL INFORMATION:
 Sequence 9, Application US/11090916 Publication No. US20050260140A1
 Matches
 Query Match
 APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
 PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR PPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
 APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
 APPLICANT: White, Carl W. APPLICANT: del Val, Greg APPLICANT: Lee, Rees
 NUMBER OF SEQ ID NOS: 15
 PRIOR APPLICATION NUMBER: 60/650,865 PRIOR FILING DATE: 2005-02-07
 PRIOR APPLICATION NUMBER: 60/556,516 PRIOR FILING DATE: 2004-03-24
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 1 WCGPCK 6
 1 WCGPCK 6
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Pred. No. 0.25;
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 บร-11-090-916-12
 US-11-090-916-10
 PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR FILING DATE: 2004-03-24
PRIOR FILING DATE: 2004-03-24
 Sequence 12, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
 SEQ ID NO 10
LENGTH: 105
 GENERAL INFORMATION:
APPLICANT: White, C
 Query Match
Best Local Similarity
 SEQ ID NO 11
 Sequence 11, Application US/11090916
Publication No. US20050260140A1
 Query Match
 Matches
 Matches
 APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
 PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
SOPTWARE: PatentIn version 3.3
 APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
 NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn versi
 FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
CURRENT FILING DATE: 2004-09,960
PRIOR APPLICATION NUMBER: 60/409,960
 PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
PRIOR APPLICATION NUMBER: 60/462,082
 ORGANISM: Rattus norvegicus
 TYPE: PRT
 Local
 31 WCGPCK 36
 31 WCGPCK
 1 Similarity 6; Conserv
 1 WCGPCK 6
 1 WCGPCK 6
 9
 PatentIn version 3.3
 del Val,
 100.0%;
ilarity 100.0%;
Conservative 0
 Conservative
 Carl W.
 100.0%; Score 47;
100.0%; Pred. No.
 0,
 ,
,
 Score 47; DB 7
Pred. No. 0.25;
 Mismatches
 Mismatches
 0.25;
 DB 7; Length 105;
 DB 7;
 Length 105
 Indels
 0,
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 Gaps
 Gaps
 0
 0
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RESULT 13
US-11-144-301A-9
 문
 S
 ; LENGTH: 105
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-106-796-13
 RESULT 12
US-11-106-796-13
 밁
 5
 US-11-090-916-12
 Sequence 13, Application US/11106796
Publication No. US20050282191A1
GENERAL INFORMATION:
APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
FILE REFERENCE: UVMO:033US
CURRENT FILING DATE: 205-04-15
PRIOR APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2004-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 13
 Sequence 9, Application US/11144301A Publication No. US20050288227A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 6; Conserv
 Best Local Similarity
Matches 6; Conserv
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 12
 Query Match
APPLICANT: Johanna, Ungerstedt
TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
FILE REFERENCE: 28517-501 UTIL
CURRENT APPLICATION NUMBER: US/11/144,301A
CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US 60/577,089
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR PRIOR PRIOR DATE: 2003-02-15
RIOR RILING DATE: 2003-02-15
NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
NUMBER: OF SEQ ID NOS: 10
 PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
 APPLICANT: Marks, Paul A. APPLICANT: Johanna, Unge
 WHIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
 PRIOR FILING DATE: 2003-04-11
 TYPE: PRT
ORGANISM: Homo sapiens
 LENGTH: 105
 31
 31
 1 WCGPCK 6
 WCGPCK 36
 WCGPCK 6
 WCGPCK 36
 100.0%; Score 47; DB 7; ilarity 100.0%; Pred. No. 0.25; Conservative 0; Mismatches
 Conservative
 100.0%; Score 47; DB 7
100.0%; Pred. No. 0.25;
 <u>,,</u>
 Mismatches
 DB 7;
 DB 7; Length 105;
 0
 <u>.</u>
 Length 105
 0;
 0,
 Gaps
 Gaps
 0
 0
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TITLE OF INVENTION: DESIGN OF THERAPEUTICS AND THERAPEUTICS
FILE REFERENCE: 17102-013001 / 1762
CURRENT APPLICATION NUMBER: US/11/032,773
CURRENT FILING DATE: 2005-01-11
PRIOR APPLICATION NUMBER: 60/536,184
PRIOR FILING DATE: 2004-01-12
PRIOR APPLICATION NUMBER: 60/536,184
PRIOR FILING DATE: 2004-03-29
INUMBER: 60/557,591
PRIOR PILING DATE: 2004-03-29
INUMBER: OF SEQ ID NOS: 958
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 943
LENGTH: 105
TYPE: PRT
ORGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank BAA04881
 RESULT 15
US-11-032-773-943
 밁
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 Ś
 Query Match
Best Local Similarity
Thes 6; Conserve
 US-11-144-301A-9
 US-11-144-301A-10
 CURRENT APPLICATION NUMBER: US/11/144,301A
CURRENT FILING DATE: 2005-06-03
PRIOR APPLICATION NUMBER: US 60/577,089
PRIOR FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR APPLICATION NUMBER: US 10/369,094
PRIOR FILING DATE: 2003-02-14
PRIOR PPLICATION NUMBER: US 60/357,383
PRIOR FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 10
 US-11-144-301A-10
 Sequence 943, Application US/11032773 Publication No. US20060018911A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 10
 SEQ ID NO 9
 Matches
 Query Match
 GENERAL INFORMATION:
 Sequence 10, Application US/11144301A Publication No. US20050288227A1
 APPLICANT: Ault-Riche, Dana APPLICANT: Levy, Ronald
 APPLICANT: Marks, Paul A. APPLICANT: Johanna, Unge
 FILE REFERENCE: 28517-501 UTIL
 APPLICANT: JOHANNA, UNGERSTECKT
TITLE OF INVENTION: USE OF THIOREDOXIN MEASUREMENTS FOR DIAGNOSTICS AND TREATMENTS
 LENGTH: 105
TYPE: PRT
ORGANISM: Homo sapiens
 TYPE: PRT
ORGANISM: Homo sapiens
DATABASE ENTRY DATE: 2002-12-25
 LENGTH:
 / Match 100.0%;
Local Similarity 100.0%;
les 6; Conservative 0
 31
 31 WCGPCK 36
 1 WCGPCK 6
 105
 1 WCGPCK 6
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 Conservative
 100.0%;
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 Score 47; DB 7
Pred. No. 0.25;
 Score 47; DB 7
Pred. No. 0.25;
Mismatches
 DB 7;
 DB 7; Length 105;
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 Length 105
 Indels
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 0;
 Gaps
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 0
 0,
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Query Match
Best Local Similarity
'Atches 6; Conserv
 문
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 ; TYPE: PRT ; ORGANISM: Mus musculus US-11-106-796-11
 밁
 S
 ; TYPE: PRT
; ORGANISM: Pseudomonas syringae
US-11-090-916-4
 US-11-106-796-11
 FILE REFERENCE: 2879-98-1

CURRENT APPLICATION NUMBER: US/11/090,916

CURRENT FILING DATE: 2005-03-24

PRIOR APPLICATION NUMBER: 60/409,960

PRIOR APPLICATION NUMBER: 60/462,082

PRIOR PILING DATE: 2003-04-11

PRIOR FILING DATE: 2003-04-11

PRIOR PILING DATE: 2003-05-10

PRIOR APPLICATION NUMBER: 60/556,516

PRIOR APPLICATION NUMBER: 60/556,516

PRIOR APPLICATION NUMBER: 60/556,516

PRIOR PILING DATE: 2004-03-24

PRIOR FILING DATE: 2004-03-24

PRIOR PILING DATE: 2004-03-24
 US-11-090-916-4
 RESULT 16
 US-11-032-773-943
 SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 11
LENGTH: 127
 SOFTWARE: PatentIn
 Sequence 11, Application US/11106796
Publication No. US20050282191A1
GENERAL INFORMATION:
 Sequence 4, Application US/11090916
Publication No. US20050260140A1
Query Match
Best Local Similarity
 Matches
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 PILE REFERENCE: UVMO:033US
CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
 APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OP INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
 APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
 NUMBER OF SEQ ID NOS: 15
 PRIOR FILING DATE:
 LENGTH: 109
 APPLICATION NUMBER: 60/650,865
 31 WCGPCK 36
 зз wcgpck зв
 1 WCGPCK 6
 1 WCGPCK 6
 Conservative
 Conservative
 version 3.3
 2005-02-07
 100.0%; Score 47; DB 7
100.0%; Pred. No. 0.26;
100.0%;
 100.0%; Score 47; I
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 <u>,</u>
Score 47; Pred. No.
 Mismatches
 Mismatches
; DB 7;
. 0.29;
 DB 7;
 DB 7; Length 105;
 0,
 0
 Length 109
 Length 127;
 Indels
 0
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 Gaps
 0
 0;
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RESULT 20
US-11-131-744-3
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 ; ORGANISM: RAT US-11-106-796-12
 ; ORGANISM: Neisseria gonorrhoeae US-10-467-657-4354
 US-11-106-796-12
 US-10-467-657-4354
 SOFTWARE: PatentIn V
SEQ ID NO 12
LENGTH: 127
TYPE: PRT
 SOPTWARE: SeqWin99, version 1.04
SEQ ID NO 4354
LENGTH: 129
 Sequence 4354, Application US/10467657 Publication No. US20050260581A1 GENERAL INFORMATION:
 Matches
 Best Local Similarity
Matches 6; Conserv
 Sequence 12, Application US/11106796 Publication No. US20050282191A1
 Query Match
 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 Matches
 APPLICANT: PIZZA Mariagrazia
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Elisabetta
 APPLICANT: SUTOVSKY, PETER
APPLICANT: MIRANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
FILE REFERENCE: UVMO:033US
 TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS FILE REFERENCE: FILE REPERENCE: US/10/467,657 CURRENT FILING DATE: 2003-08-11 PRIOR APPLICATION NUMBER: GB-0103424.8 PRIOR APPLICATION NUMBER: GB-0103424.8 PRIOR FILING DATE: 2001-02-12 NUMBER OF SEQ ID NOS: 9218
 PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
 CURRENT APPLICATION NUMBER: US/11/106,796
CURRENT FILING DATE: 2005-04-15
 APPLICANT: CHIRON SPA
APPLICANT: FONTANA Maria Rita
 TYPE: PRT
 31
 31
 52 WCGPCK 57
 1 WCGPCK 6
 1 WCGPCK 6
 6;
 6,
 9
 100.0%;
nilarity 100.0%;
Conservative 0
 Conservative
 100.0%; Score 47; DB 7; Length 127; ilarity 100.0%; Pred. No. 0.29; Conservative 0; Mismatches 0; Indels
 Ver. 2.1
 0;
 0,
 Score 47; DB 6;
Pred. No. 0.29;
0; Mismatches 0
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 Indels
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 Gaps
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 0
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; Sequence 3, Application US/11131744

Publication No. US20060023808A1

APPLICANT: HAJIVANDI, MAHBOD R.

SADEGHI, MEHRNOOSH

ROBERT M.

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멍
 US-11-131-744-6
 US-11-131-744-3
 SOFTWARE: PatentIn
SEQ ID NO 6
 Sequence 6, Application US/11131744 Publication No. US20060023808A1
 Matches
 Query Match
 SEQ ID NO 3
 SENERAL INFORMATION
 PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR PRIOR PRIOR NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
PRIOR FILING DATE: 2004-05-17
 PRIOR APPLICATION NUMBER: 60/572,290
PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
PRIOR FILING DATE: 2004-05-17
 CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
 TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR CALIBRATION IN MASS TITLE OF INVENTION: SPECTROMETRY FILE REFERENCE: NVX-28 US
 APPLICANT:
 NUMBER OF SEQ ID NOS: 9
 PRIOR APPLICATION NUMBER: 60/621,686
PRIOR FILING DATE: 2004-10-26
 CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
 APPLICANT: LIANG, XIQUAN
APPLICANT: LETTE, JOHN F.
TITLE OF INVENTION: COMPOSITIONS,
TITLE OF INVENTION: SPECTROMETRY
 PRIOR FILING DATE:
 APPLICANT:
 APPLICANT: HAJIVANDI, MAHBOD R. APPLICANT: POPE, ROBERT M.
 SOFTWARE: PatentIn Ver. 3.3
 NUMBER OF SEQ
 FILE REFERENCE: NVX-28 US
 FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: protein sequence
 ORGANISM: Artificial Sequence
DRGANISM: Artificial Sequence
 ENGTH:
 ENGTH: 177
 Local Similarity nes 6; Conserv
 FILING DATE: 2004-05-17
APPLICATION NUMBER: 60/621,685
 APPLICATION NUMBER: 60/621,686
 APPLICATION NUMBER: 60/621,685 FILING DATE: 2004-10-26
 FILING DATE: 2004-10-26
 31 WCGPCK 36
 273
 1 WCGPCK 6
 SADEGHI, MEHRNOOSH
 LEITE, JOHN F.
 ID NOS:
 Conservative
 Ver. 3.3
 2004-10-26
 100.0%; Score 47; 100.0%; Pred. No.
 0
 KITS, AND METHODS FOR CALIBRATION IN MASS
 В
 7;
 Length 177;
 Indels
 0;
 Gaps
 0
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RESULT 22
US-11-131-744-7
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 ; PEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: protein sequence
US-11-131-744-6
 US-11-090-916-6
Sequence 6, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
APPLICANT: White, Carl W.
APPLICANT: Lee, Rees
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
PILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
 Query Match
Best Local Similarity
 SOFTWARE: PatentIn Ver. 3.3 SEQ ID NO 7
 Sequence 7, Application US/11131744 Publication No. US20060023808A1
 Matches
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 Query Match
 GENERAL INFORMATION
 Best Local Similarity 100.
Matches 6; Conservative
 PRIOR FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/669,373
PRIOR FILING DATE: 2005-04-08
PRIOR APPLICATION NUMBER: 60/572,215
PRIOR FILING DATE: 2004-05-17
 CURRENT APPLICATION NUMBER: US/11/131,744
CURRENT FILING DATE: 2005-05-17
PRIOR APPLICATION NUMBER: 60/572,290
 PRIOR APPLICATION NUMBER: 60/621,686 PRIOR FILING DATE: 2004-10-26 NUMBER OF SEQ ID NOS: 9
 APPLICANT: HAJIVANDI, MAHBOD R. APPLICANT: POPE, ROBERT M.
 TITLE OF INVENTION: COMPOSITIONS, TITLE OF INVENTION: SPECTROMETRY FILE REFERENCE: NVX-28 US
 APPLICANT:
 APPLICANT:
 OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: protein sequence
 TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
 APPLICANT:
 ENGTH: 440
 APPLICATION NUMBER: 60/621,685 FILING DATE: 2004-10-26
 32 WCGPCK 37
 31 WCGPCK 36
 1 WCGPCK 6
 1 WCGPCK 6
 6
 LIANG, XIQUAN
LEITE, JOHN F.
 LIANG,
 SADEGHI, MEHRNOOSH
 Conservative
 100.0%;
 100.0%; Score 47; DB 7
100.0%; Pred. No. 0.65;
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 Score 47;
Pred. No.
 Mismatches
 KITS, AND METHODS FOR CALIBRATION IN MASS
 DB 7; Length 440
 DB 7;
 0
 Length 273
 Indels
 0;
 0
 Gaps
 Gaps
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RESULT 24
US-11-090-916-5
 S
 ; LENGTH: 103
; TYPE: PRT
; ORGANIZM: Listeria monocytogenes
US-11-090-916-6
 RESULT 25
US-11-098-686-10612
 밁
 S
 US-11-090-916-5
 PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR FILING DATE: 2003-09-10
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
 Sequence 10612, Application US/11098686
publication No. US20060024696A1
GENERAL INFORMATION:
APPLICANT: Kapur, Vivek and Gebhart, Connie J.
TITLE OF INVENTION: NUCLEIC ACID AND POLYPEPTIDE SEQUENCES
TITLE OF INVENTION: FROM LAWSONIA INTRACELLULARIS AND METHODS OF USING
 NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin version 3.3
SEQ ID NO 5
LENGTH: 104
 Sequence 5, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 5; Conserv
 Query Match
Best Local Similarity
Matches 5; Conserv
 SOFTWARE: PatentIn version 3.3 SEQ ID NO 6
 APPLICANT: White, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
 PRIOR APPLICATION NUMBER: 60/556,516
PRIOR PILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
 PRIOR PRIOR
 PRIOR APPLICATION NUMBER: 60/409,960 PRIOR FILING DATE: 2002-09-10
 PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
 PRIOR FILING DATE: 2005-02-07
FILE REFERENCE:
 TYPE: PRT ORGANISM: Porphyromonas gingivalis
 APPLICATION NUMBER: 10/660,118 FILING DATE: 2003-09-10
 27 WCGPCR 32
 28 WCGPCR 33
 1 WCGPCK 6
 1 WCGPCK 6
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 93.6%;
 93.6%;
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 Score 44; DB 7;
Pred. No. 0.74;
 Score 44; DB 7
Pred. No. 0.74;
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 DB 7;
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 0; Indels
 Length 104;
 Length 103;
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 0;
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Sequence 12, Application US/11195459
Publication No. US20050278803A1
GENERAL INFORMATION:
APPLICANT: Sewalt, Vincent
APPLICANT: Hastings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
 SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10612
LENCTH: 107
TYPE: PRT
ORGANISM: Lawsonia intracellularis
US-11-098-686-10612
 US-11-195-459-16
US-11-195-459-16
; Sequence 16, Application US/11195459
; Publication No. US20050278803A1
; GENERAL INFORMATION:
; APPLICANT: Sewalt, Vincent
; APPLICANT: Hastings, Craig
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 US-11-195-459-12
 US-11-195-459-12
 Best Loc
Matches
 NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn version 3.0
SEQ ID NO 12
LENGTH: 122
 Query Match
Best Local S
 APPLICANT:
 Query Match
 Matches
 APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PRO
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
CURRENT FILING DATE: 2005-08-02
CURRENT APPLICATION NUMBER: US/10/005,429
 CURRENT APPLICATION NUMBER: US/11/098,686
CURRENT FILLING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US03/31318
PRIOR FILING DATE: 2003-10-01
PRIOR APPLICATION NUMBER: US 60/416,395
PRIOR FILING DATE: 2002-10-04
NUMBER OF SEQ ID NOS: 11433
 APPLICANT:
 PRIOR FILING DATE: 2001-12-03
PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
 APPLICANT:
 APPLICANT:
 TYPE: PRT
ORGANISM: Zea mays
 Local Similarity
 Local Similarity
les 5; Conserv
 ç
 39 WCGPCR 44
 29 WCGPCR
T: Jung, Rudolf
T: Everard, John
T: Allen, Stephen
INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS
 1 WCGPCK 6
 1 WCGPCK 6
 Hastings, Craig
Meeley, Robert
Hantke, Sabine
Jung, Rudolf
 Conservative
 Conservative
 34
 93.6%;
 93.6%; Score 44; DB 7;
83.3%; Pred. No. 0.76;
 Score 44; DB 7
Pred. No. 0.82;
 Mismatches
 Mismatches
 DB 7;
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 0;
 Length 122;
 Length 107;
 Indels
 <u>,</u>
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 Gaps
 Gaps
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 0;
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PR

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CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin version 3.0
SEQ ID NO 16
LENGTH: 122
TYPE: PRT
ORGANISM: Hordeum vulgare
 RESULT 29
US-11-195-459-10
 ; FEATURE:
, NAME/KEY: misc feature
, OTHER INFORMATION: "n" at position 9,
, OTHER INFORMATION: or t
US-11-195-459-19
 APPLICANT: Hastings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
APPLICANT: Jung, Rudolf
APPLICANT: Everard
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 RESULT 28
US-11-195-459-19
 밁
 US-11-195-459-16
 Sequence 10, Application US/11195459
Publication No. US20050278803A1
GENERAL INFORMATION:
APPLICANT: Sewalt, Vincent
APPLICANT: Hastings, Craig
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 19
LENGTH: 122
 Matches
 Query Match
Best Local Similarity
 Query Match
Best Local
 APPLICANT: EVERAL, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
NUMBER OF SEQ ID NOS: 25
 GENERAL INFORMATION:
 Sequence 19, Application US/11195459 Publication No. US20050278803A1
 Matches
 APPLICANT: Sewalt, Vincent
 FILE REFERENCE: 5718-119 (035718/241421)
 APPLICANT:
 ORGANISM: Zea mays
 TYPE: PRT
 39 WCGPCR 44
 45
 1 WCGPCK 6
 1 WCGPCK 6
 <u>ب</u>
 Similarity
 WCGPCR 50
Hastings, Crai
Meeley, Robert
 Conservative
 Conservative
 93.6%;
 93.6%;

 Mismatches

 Score 44; DB 7;
Pred. No. 0.82;
1; Mismatches
 Score 44; DB 7
Pred. No. 0.82;
 493, 537,
 DB 7;
 0;
 Length 122
 548,
 Length 122
 Indels
 581,
 and 584 can be
 0
 0,
 Gaps
 Gaps
 0
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 0;
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 RESULT 31
US-11-195-459-8
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APPLICANT: Maite, Carl W.
APPLICANT: del Val, Greg
APPLICANT: Lee, Rees
ITILE OF INVENTION:
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR REPLICATION NUMBER: 60/409,960
PRIOR REPLICATION NUMBER: 60/462,082
PRIOR REPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/550,865
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR APPLICATION NUMBER: 60/650,865
밁
 밁
 ; ORGANISM: Zea mays US-11-195-459-10
 ; ORGANISM: Arabidopsis thaliana US-11-090-916-13
 US-11-090-916-13
 SOFTWARE: PatentIn version 3.3
SEQ ID NO 13
LENGTH: 134
 SOFTWARE: PatentIn version 3.0
SEQ ID NO 10
LENGTH: 123
TYPE: PRT
 GENERAL INFORMATION:
 Matches
 Query Match
Best Local
 Sequence 13, Ap
Publication No.
 Query Match
Best Local Similarity
 APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-01
PRIOR FILING DATE: 2001-12-01
PRIOR FILING DATE: 2001-12-01
PRIOR FILING DATE: 2000-12-01
 Matches
 NUMBER OF SEQ ID NOS: 15
 TYPE: PRT
 Local Similarity
 11111:
40 WCGPCR 45
59 WCGPCR
 1 WCGPCK 6
 1 WCGPCK 6
 5
 Application US/11090916
No. US20050260140A1
 Hantke, Sabine
 Conservative
 Conservative
64
 93.6%;
 93.6%; Score 44; DB 7;
83.3%; Pred. No. 0.88;
 Score 44; I
Pred. No. 0
 Mismatches
 Mismatches
 0
 0
 Length 134;
 Length 123
 Indels
 0
 0
 Gaps
 Gaps
 0
 0
 ဝှ
 PR
```

Sequence 8, Application US/11195459
Publication No. US20050278803A1

GENERAL INFORMATION:

```
APPLICANT: Sewall,
APPLICANT: Hastings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Heathe, Sabine
APPLICANT: Heathe, Sabine
APPLICANT: Heathe, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PROTECTION ENGRAPHED IN NUMBER: US/11/195,459
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR APPLICATION NUMBER: 00/250,703
PRIOR APPLICATION NUMBER: 00/250,703
PRIOR APPLICATION NUMBER: 00/250,703
RESULT 33
US-11-090-916-15
 RESULT 32
US-11-090-916-14
 Query Match
Best Local Similarity
Tatches 5; Conserv
 문
 밁
 ; TYPE: PRT; ORGANISM: Zea mays US-11-195-459-8
 Ś
 US-11-090-916-14
 Query Match
Best Local Similarity
Matches 5; Conserv
 NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn version 3.3
SEQ ID NO 14
LENGTH: 167
 Sequence 14, Application US/11090916
Publication No. US20050260140A1
GENERAL INFORMATION:
 TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof FILE REFERENCE: 2879-98-1
CURRENT APPLICATION NUMBER: US/11/090,916
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR FILING DATE: 2002-09-10
PRIOR FILING DATE: 2002-09-11
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2003-04-11
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2004-03-24
PRIOR APPLICATION NUMBER: 60/650,865
 APPLICANT: White, Carl W. APPLICANT: del Val, Greg
 PRIOR FILING DATE:
 TYPE: PRT
ORGANISM: Zea mays
 LENGTH: 134
 S4 WCGPCR 59
 88 WCGPCR 93
 1 WCGPCK 6
 1 WCGPCK 6
 Conservative
 Conservative
 Rees
 Carl W.
 2005-02-07
 93.6%;
 93.6%;

 Mismatches

 Score 44; DB 7;
Pred. No. 0.88;
 Score 44; D
Pred. No. 1;
 Mismatches
 0; Indels
 0;
 Length 134;
 Indels
 0
 ,
,
 Gaps
 Gaps
 0;
 0
```

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Sequence 8494, Application US/10467657
Publication No. US2005280581A1
GENERAL INFORMATION:
APPLICANT: CHIRON SpA
APPLICANT: FONTANA Maria Rita
APPLICANT: MASIGNANI Vega
APPLICANT: MASIGNANI Vega
APPLICANT: MONACI Elisabetta
 밁
 Ş
 ; ORGANISM: Oryza sativa
US-11-090-916-15
 밁
 US-10-467-657-8494
 ; ORGANISM: Neisseria gonorrhoeae
US-10-467-657-8494
 Query Match
Best Local S
Matches 5
 APPLICANT: Clee, Rees
APPLICANT: Lee, Rees
TITLE OF INVENTION: Mucolytic and Anti-Elastase Compounds and Methods of Use Thereof
FILE REFERENCE: 2879-98-1
CURRENT FILING DATE: 2005-03-24
PRIOR APPLICATION NUMBER: 60/409,960
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR FILING DATE: 2002-09-10
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 60/462,082
PRIOR APPLICATION NUMBER: 10/660,118
PRIOR APPLICATION NUMBER: 10/650,118
PRIOR APPLICATION NUMBER: 60/556,516
PRIOR FILING DATE: 2003-09-10
PRIOR PRIOR PRICE 2003-03-24
PRIOR PRICE 2003-03-24
PRIOR PRICE 2003-03-24
PRIOR PRICE 2003-03-24
 CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
PRIOR APPLICATION NUMBER: GB-0103424.8
PRIOR FILING DATE: 2001-02-12
RUMBER OF SEQ ID NOS: 9218
SOFTWARE: SeqWin99, version 1.04
SEQ ID NO 8494
LENGTH: 209
 SEQ ID NO 15
 Sequence 15, Application US/11090916 Publication No. US20050260140A1
 GENERAL INFORMATION:
 Matches
 Query Match
Best Local Similarity
 PRIOR APPLICATION NUMBER: 60/650,865
PRIOR FILING DATE: 2005-02-07
NUMBER OF SEQ ID NOS: 15
 TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS FILE REFERENCE:
 LENGTH: 17
TYPE: PRT
 SOFTWARE:
 TYPE: PRT
 Local Similarity tes 5; Conserv
101 WCGPCR 106
 94 WCGPCR
 172
 1 WCGPCK 6
 1 WCGPCK 6
 ۍ
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 PatentIn version 3.3
 Conservative
 Conservative
 99
 Carl W.
 93.6%;
 93.6%; Score 44; DB 6; 83.3%; Pred. No. 1.2;
 Score 44; DB 7;
Pred. No. 1;
1; Mismatches
 0
 Length 209;
 Length 172;
 Indels
 0
 o,
 Gaps
 Gaps
 0
 0
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RESULT 35 US-10-995-561-703 ; Sequence 703, Application US/10995561

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CURRENT FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: 60/049911
PRIOR APPLICATION NUMBER: 60/05974
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059117
PRIOR PILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR APPLICATION NUMBER: 60/059184
 밁
 RESULT 36
 ; ORGANISM: Homo sapiens
US-10-995-561-703
 US-10-131-826A-426
 NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FRSTSEQ for Windows Version 4.0
SEQ ID NO 703
LENGTH: 126
TYPE: PRT
 Sequence 426, Application US/10131826A Publication No. US20050245730A1
 Matches
 Query Match
Best Local Similarity
 GENERAL INFORMATION:
 Publication No. US20050272054A1
 APPLICANT:
 APPLICANT:
 APPLICANT: CARGIIL, Michele et al.
TITLE OF INVENTION: GENETIC COULAR DISORDERS AND DRUG RESPONSE, METHODS
TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CL001559
CURRENT PEPLICATION NUMBER: US/10/995,561
CURRENT FILING DATE: 2004-11-24
 APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C128
 APPLICANT: Baker, Kevin P.
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICATION NUMBER: 60/059263
PILING DATE: 1997-09-19
APPLICATION NUMBER: 60/059588
 APPLICATION NUMBER: 60/059352
 62 WCGPCQ 67
 1 WCGPCK 6
 <u>ب</u>
 Godowski, Paul J. Gurney, Austin L.
 Goddard, Audrey
 Gerritsen, Mary E
 Wood, William
 Watanabe, Colin K
 Tumas,Daniel
 Stewart, Timothy A
 Smith, Victoria
 Sherwood, Steven
 Filvaroff,Ellen
 DeForge, Laura
 Beresini,Maureen
 Conservative
 1997-09-18
 91.5%;
83.3%;
 Score 43;
Pred. No.
 Mismatches
 6,
 0;
 Length 126;
 Indels
 0,
 Gaps
 얹
 0
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 RESULT 38
US-10-821-234-860
 밁
 RESULT 37
US-10-467-962B-6
 US-10-131-826A-426
 US-10-467-962B-6
 SEQ ID NO 6
LENGTH: 183
TYPE: PRT
 SOFTWARE: pt
SEQ ID NO 860
LENGTH: 176
 Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 426
LENGTH: 747
TYPE: PRT
ORGANISM: Homo Sapien
 Sequence 860, Application US/10821234 Publication No. US20050255114A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 Matches
 Matches
 GENERAL INFORMATION:
 Sequence 6, Application US/1046 Publication No. US20050246784A1
 Query Match
 APPLICANT: Plesch, Gunnar
APPLICANT: Blau, Astrid
APPLICANT: Blau, Astrid
APPLICANT: Daeschner, Klaus
APPLICANT: Klein, Mathieu
 Best Local Similarity
 FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
CURRENT FILING DATE: US 60/462,047
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
 FILE REFERENCE: 2000_857
CURRENT APPLICATION NUMBER: US/10/467,962B
CURRENT FILING DATE: 2003-08-14
 PRIOR APPLICATION NUMBER: PCT/EP02/01466
PRIOR FILING DATE: 2002-02-13
NUMBER OF SEQ ID NOS: 109
 TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
 APPLICANT:
 APPLICANT:
 APPLICANT: Labat, Ivan
 SOFTWARE: PatentIn Vers. 2.0
 TITLE OF INVENTION: Identification of Herbicidally Active Substances
TYPE: PRT
ORGANISM: Homo sapiens
 ORGANISM: Arabidopsis thaliana
 105 WCGPC 109
 653 WCGPCQ 658
 5;
 1 WCGPC 5
 1 WCGPCK 6
 5
 Application US/10467962B
 Andarmani, Su
Tang, Y. Tom
 Stache-Crain, Birgit
 _SEQ_genes Version 1.0
 Conservative
 Conservative
 100.0%;
 91.5%;
83.3%;
 Susan
 89.4%; Score 42;
100.0%; Pred. No.
 0;
 1; Mismatches
 Score 43; DB 6;
Pred. No. 3.8;
 Mismatches
 DB 6;
 Length 183;
 Length 747;
 Indels
 ç
 0
 Gaps
 0;
 0,
```

Query Match

80.9%;

Score 38;

DB 6;

Length 176;

83.3%; Pred. No.

9

```
Sequence 114, Application US/10878556A
Publication No. US20050266399A1
GENERAL INFORMATION:
APPLICANT: Hoffmann La-Roche Inc.
TITLE OF INVENTION: HCV regulated protein expression
FILE REFERENCE: 21762
CURRENT APPLICATION NUMBER: US/10/878,556A
CURRENT FILING DATE: 2004-06-28
NUMBER OF SEQ ID NOS: 199
SOFTWARE: Patentin version 3.1
SEQ ID NO 114
LENGTH: 324
 APPLICANT: Zhou, Ping
APPLICANT: Cao, Yi-Cheng
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
FILE REFERENCE: 785CIP4CN
CURRENT APPLICATION NUMBER: US/11/000,463
CURRENT FILING DATE: 2004-11-29
PRIOR APPLICATION NUMBER: 10/291,265
PRIOR APPLICATION NUMBER: 10/291,265
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 09/922,279
PRIOR APPLICATION NUMBER: 09/922,279
PRIOR APPLICATION NUMBER: 09/921,404
PRIOR APPLICATION NUMBER: 09/491,404
PRIOR APPLICATION NUMBER: 09/617,746
PRIOR APPLICATION NUMBER: 09/631,451
PRIOR APPLICATION NUMBER: 09/631,451
PRIOR APPLICATION NUMBER: 09/631,451
PRIOR APPLICATION NUMBER: 09/631,451
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 RESULT 40
US-10-878-556A-114
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 US-11-000-463-344
 SOFTWARE: Fas
SEQ ID NO 344
LENGTH: 269
 Query Match
Best Local Similarity
Matches 5; Conserv
 Best Local Similarity Matches 5; Conserv
 Sequence 344, Publication No.
 GENERAL INFORMATION
 APPLICANT:
APPLICANT:
APPLICANT:
 NUMBER OF SEQ ID NOS: 944
COSTWARE: FastSEQ for Windows Version 3.0
 APPLICANT:
 APPLICANT: Tang, APPLICANT: Liu,
 PRIOR FILING DATE:
 APPLICANT:
 TYPE: PRT
ORGANISM: Homo sapiens
 APPLICATION NUMBER: 09/633,870
 53 WCGHCK 58
 93 WCGHCK 98
 1 WCGPCK 6
 1 WCGPCK 6
 Qian, Xiaoho
Wang, Zhiwei
 Wehrman,
 Zhang, Jie
Zhou, Ping
 Liu, Chenghua
Asundi, Vinod
 Application US/11000463
DS US20050266423A1
 Conservative
 Conservative
 Rui-hong
 Xiaohong
 Tom
 2000-09-15
 80.9%;
 в.
 Score 38; DB Pred. No. 12; 0; Mismatches
 0
 <u>,</u>
 Mismatches
 7;
 1;
 Length 269;
 Indels
 0;
 0,
 Gaps
 Gaps
 0
 0
```

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Sequence 70, Application US/10523503

Publication No. US20060037102A1

GENERAL INFORMATION:
APPLICANT: BASE PLANT SCIENCE GMBH
TITLE OF INVENTION: SUGAR AND LIPID METABOLISM REGULATORS IN PLANTS IV
FILE REFERENCE: 16313-0236

CURRENT APPLICATION NUMBER: US/10/523,503

CURRENT FILING DATE: 2005-02-02
PRIOR APPLICATION NUMBER: US 60/400,803
PRIOR FILING DATE: 2002-08-02
INUMBER OF SEQ ID NOS: 163
SOFTWARE: Patentin version 3.2

SEQ ID NO 70
LENGTH: 361
TYPE: PRT
 Sequence 1, Application US/1087856A
Publication No. US20050266399A1
GENERAL INFORMATION:
APPLICANT: Hoffmann La-Roche Inc.
TITLE OF INVENTION: HCV regulated protein expression
FILE REFERENCE: 21762
CURRENT APPLICATION NUMBER: US/10/878,556A
CURRENT FILING DATE: 2004-06-28
NUMBER OF SEQ ID NOS: 199
SOPTWARE: PatentIn version 3.1
SEQ ID NO 1
LENGTH: 414
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 US-10-878-556A-114
 US-10-878-556A-1
 US-10-878-556A-1
 ; ORGANISM: Arabidopsis thaliana US-10-523-503-70
 US-10-523-503-70
 RESULT 41
 Best Local Similarity
Matches 5; Conserv
Query Match
Best Local Similarity
Matches 5; Conserv
 Query Match
 Matches
 Query Match
Best Local Similarity
 NAME/KEY: MISC FEATURE
LOCATION: (307)..(307)
OTHER INFORMATION: Xaa is an undetermined amino acid residue
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: hsugp:006101-3-0
DATABASE ENTRY DATE: 2003-02-16
 ORGANISM: Homo sapians
PUBLICATION INPORMATION:
DATABASE ACCESSION NUMBER: tr hum/q9bvh9
DATABASE ENTRY DATE: 2001-06-01
 ORGANISM: Homo sapiens FEATURE:
 TYPE: PRT
 108 WCGHCK 113
 51
 1 WCGPCK 6
 1 WCGPCK 6
 5
 WCGHCK 56
 Conservative
 Conservative
 Conservative
 80.9%;
 80.9%;
 80.9%; Score 38; 83.3%; Pred. No.
Score 38; DB 6; Pred. No. 16; 0; Mismatches
 0;
 .
 Score 38;
Pred. No.
 Mismatches
 Mismatches
 DB
13;
 DB
14;
 6,
 6,
 ۲.
 1;
 Length 414;
 Length 324;
 Length 361;
 Indels
 Indels
 0,
 0
 0;
 Gaps
 Gaps
 Gaps
 0
 0
 <u>..</u>
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WCGPCK 6

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PRIOR APPLICATION UNMBER US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
SOTTWARE: pt SEQ_genes Version 1.0
SEQ ID NO 1287
LENGTH: 505
TYPE: PRT
ORCANISM: Homo sapiens
US-10-821-234-1287
 PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
SOPTWARE: pt SEQ_genes Version 1.0
SEQ ID NO 1282
SEQ ID NO 1282
LENGTH: 440
TYPE: PRT
ORGANISM: Homo sapiens
US-10-821-234-1282
RESULT 45
US-10-821-234-1409
; Sequence 1409, Application US/10821234
 В
 Query Match
Best Local Similarity
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 밁
 US-10-821-234-1287
 US-10-821-234-1282
 Sequence 1282, Application US/10821234
Publication No. US20050255114A1
GENERAL INFORMATION:
APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit
 Sequence 1287, Application US/10821234 Publication No. US20050255114A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
Matches 5; Conserv
 APPLICANT: Andarmani, Susan
APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
CURRENT FILING DATE: 2004-04-07
 FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT FILING DATE: 2004-04-07
 APPLICANT: Andarmani, Susan
APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
 APPLICANT: Labat, Ivan
APPLICANT: Stache-Crai
 189 WCGHCK 194
 236
 1 WCGPCK 6
 1 WCGPCK 6
 WCGHCK 61
 WCGHCK 241
 Stache-Crain,
 Conservative
 Conservative
 80.9%;
 80.9%;
 Birgit
 0
 Score 38;
Pred. No.
 Score 38; DB
Pred. No. 16;
 Mismatches
 DB 6;
18;
 ٠<u>.</u>
 Length 505;
 Length 440
 0;
 0;
 0
```

```
Sequence 976, Application.
Sequence 976, Application.
Publication No. US20060034853A1
GENERAL INFORMATION:
APPLICANT: YUEN, KWOK YUNG
APPLICANT: WOO, CHIU YAT PATRICK
APPLICANT: LAU, KAR PUI SUSANNA
APPLICANT: LAU, KAR PUI SUSANNA
APPLICANT: LAU, WAR PUI SUSANNA
APPLICANT: LAU, WAR PUI SUSANNA
 RESULT 47
US-11-129-741-976
 밁
 RESULT 46
US-10-895-064-976
 밁
 ; ORGANISM: Homo sapiens US-10-821-234-1409
 US-10-895-064-976
 APPLICANT: CHAN, KNOK HUNG
APPLICANT: LEUNG, FREDERICK C.
TITLE OF INVENTION: A NOVEL HUMAN VIRUS CAUSING RESPIRATORY TRACT INFECTION AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: V0690.0031
CURRENT APPLICATION NUMBER: US/10/895,064
CURRENT FILING DATE: 2004-07-21
NUMBER OF SEQ ID NOS: 2918
SOFTWARE: Patentin version 3.2
SEQ ID NO 976
LENGTH: 32
TYPE: PRT
CRANISM: Corononavirus-HKU1
 Query Match
Best Local S
Matches 5
 SOFTWARE: pt_SEQ_genes Version 1.0 SEQ ID NO 1409
 Query Match
 APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit
APPLICANT: Andarmani, Susan
APPLICANT: Tang, Y. Tom
APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
FILE REFERENCE: 821A
CURRENT EPLICATION NUMBER: US/10/821,234
CURRENT EPLING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR APPLICATION NUMBER: US 60/462,047
NUMBER OF SEQ ID NOS: 1704
NUMBER OF SEQ ID NOS: 1704
 Matches
 Publication No. US20050255114A1
APPLICANT:
APPLICANT:
APPLICANT:
 Sequence 976, A Publication No.
 Best Local Similarity
 APPLICANT: PEIRIS, JOSEPH S.M. APPLICANT: YUEN, KWOK YUNG APPLICANT: POON, LIT MAN
 APPLICANT:
 TYPE: PRT
 ENGTH: 645
 Local Similarity
nes 5; Conserv
 INFORMATION:
 90 WCGHCK 95
 છ
 1 WCGPCK 6
 1 WCGPCK 6
 4; Conservative
 Application US/11129741
 Application US/10895064
 Conservative
 US20060018923A1
 76.6%;
 Score 36; DB Pred. No. 6.1;
 Score 38; DE Pred. No. 21;
 Mismatches
 DB 6;
 6
 Length 645;
 0
 0
 Gaps
 0
 0,
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RESULT 49
US-10-689-742-19
 밁
 S
 밁
 ; TYPE: PRT
; ORGANISM: Corononavirus-HKU1
US-11-129-741-976
 US-11-129-741-3972
 RESULT 48
 US-11-129-741-3972
 PRIOR APPLICATION NUMBER: 10/85,064
PRIOR FILING DATE: 2004-07-21
NUMBER OF SEQ ID NOS: 4257
SOFTWARE: PATENTIN VERSION 3.3
SEQ ID NO 3972
LENGTH: 32
 CURRENT APPLICATION NUMBER: US/11/129,741
CURRENT FILING DATE: 2005-05-16
PRIOR APPLICATION NUMBER: 10/895,064
PRIOR FILING DATE: 2004-07-21
NUMBER OF SEQ ID NOS: 4257
SOFTWARE: PatentIn version 3.3
SEQ ID NO 976
 Sequence 19, Application US/10689742 Publication No. US20050250180A1 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 Matches
 Query Match
 GENERAL INFORMATION:
 Sequence 3972, Application US/11129741 Publication No. US20060034853A1
 Matches
APPLICANT: Jacobs, Kenneth
APPLICANT: McCoy, John M
APPLICANT: LaVallie, Edward R
APPLICANT: Racie, Lisa A
APPLICANT: Evans, Cheryl
 APPLICANT: GUAN, YI
TITLE OF INVENTION: A NOVEL HUMAN VIRUS CAUSING RESPIRATORY TRACT
TITLE OF INVENTION: INFECTION AND USES THEREOF
FILE REFERENCE: V0690.0044
CURRENT APPLICATION NUMBER: US/11/129,741
CURRENT FILING DATE: 2005-05-16
CURRENT FILING DATE: 2005-05-16
 APPLICANT: YUEN, KWOK YUNG
APPLICANT: WOO, CHIU YAT PATRICK
APPLICANT: LAU, KAR PUI SUSANNA
 TITLE OF INVENTION: A NOVEL HUMAN VIRUS CAUSING RESPIRATORY TRACT
TITLE OF INVENTION: INFECTION AND USES THEREOF
FILE REFERENCE: V0690.0044
 LENGTH: 32
TYPE: PRT
ORGANISM: Corononavirus-HKU1
 LENGTH: 32
 PPLICANT:
 Local Similarity
 1 WCGPCK 6
 9
 1 WCGPCK 6
 9 WCGSCR 14
 4.
 4;
 WCGSCR 14
 PEIRIS, UN
 POON, LIT MAN
 PEIRIS,
 Conservative
 Conservative
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 KWOK HUNG
 JOSEPH S.M
 JOSEPH S.M.
 76.6%;
 76.6%;
66.7%;
 Score 36;
Pred. No.
 Score 36; DB 7; Length 32; Pred. No. 6.1;
 Mismatches
 Mismatches
 DB 7;
 1; Indels
 Length 32;
 0,
 0;
 Gaps
 0
 0;
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RESULT 50
US-11-106-796-10
 문
 밁
 8
 ; ORGANISM: Homo sapiens
US-10-689-742-19
 ; APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POI
FILE REFERENCE: 00766.00091.10
CURRENT APPLICATION NUMBER: US/10/689,742
CURRENT FILING DATE: 2005-10-22
PRIOR APPLICATION NUMBER: 09/746,783
PRIOR FILING DATE: 2000-12-21
NUMBER OF SEQ ID NOS: 231
NUMBER OF SEQ ID NOS: 231
APPLICANT: Hastings, Craig
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
APPLICANT: Jung, Rudolf
APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOI
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
 RESULT 51
US-11-195-459-2
 ORGANISM: Homo sapiens US-11-106-796-10
 ; SEQ ID NO 10
; LENGTH: 127
; TYPE: PRT
 Sequence 2, Application US/11195459
Publication No. US20050278803A1
GENERAL INFORMATION:
 Sequence 10, Application US/11106796 Publication No. US20050282191A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.2 SEQ ID NO 19
 Matches
 Query Match
Best Local S
 Matches
 Query Match
Best Local Similarity
 APPLICANT: SUTOVSKY, BETER
APPLICANT: MITANDA-VIZUETE, ANTONIO
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR EVALUATION OF FERTILITY
FILE REFERENCE: UVMO:033US
CURRENT FILING DATE: 2005-04-15
CURRENT FILING DATE: 2005-04-15
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR APPLICATION NUMBER: 60/562,526
PRIOR FILING DATE: 2004-04-15
NUMBER OF SEQ ID NOS: 17
 APPLICANT: Sewalt, Vincent
APPLICANT: Hastings, Crain
APPLICANT: Meeley, Robert
APPLICANT: Hantke, Sabine
 LENGTH: 107
TYPE: PRT
 Local Similarity 100.0%;
nes 5; Conservative 0;
 32 CGPCK 36
 32 CGPCK
 2 CGPCK 6
 2 CGPCK 6
 'n
 Conservative
 76.6%;
 76.6%; Score 36; DB 7; 100.0%; Pred. No. 15; ive 0; Mismatches
 0;
 Score 36; DB 6;
Pred. No. 13;
 Mismatches
 POLYNUCLEOTIDES ENCODING THEM
 FOR ALTERING THE DISULFIDE STATUS OF
 Length 127;
 Length 107;
 <u>,</u>
 0
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 0
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RESULT 53
US-11-072-512-2145
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 RESULT 52
US-11-195-459-4
 ; LENGTH: 128
; TYPE: PRT
; ORGANISM: Zea mays
US-11-195-459-2
 밁
 US-11-195-459-4
 Best Loc
Matches
 PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn version 3.0
SEQ ID NO 2
 Sequence 2145, Application US/11072512 Publication No. US20060029945A1 GENERAL INFORMATION:
 SOFTWARE: PatentIn version 3.0 SEQ ID NO 4 LENGTH: 128
 Query Match
Best Local Similarity
 Sequence 4, App. Publication No.
 Matches
 APPLICANT: Jung, Rudolf
APPLICANT: Everard, John
APPLICANT: Everard, John
APPLICANT: Allen, Stephen
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ALTERING THE DISULFIDE STATUS OF PRC
FILE REFERENCE: 5718-119 (035718/241421)
CURRENT APPLICATION NUMBER: US/11/195,459
CURRENT FILING DATE: 2005-08-02
PRIOR APPLICATION NUMBER: US/10/005,429
PRIOR FILING DATE: 2001-12-03
PRIOR FILING DATE: 2001-12-03
PRIOR APPLICATION NUMBER: 60/250,703
PRIOR FILING DATE: 2000-12-01
NUMBER OF SEQ ID NOS: 25
 GENERAL INFORMATION:
 Query Match
Best Local Similarity
 APPLICANT:
 APPLICANT:
 APPLICANT:
 APPLICANT: Sewalt, Vincent APPLICANT: Hastings, Crai
 APPLICANT:
 APPLICANT:
 APPLICANT
 PPLICANT
 TYPE: PRT
ORGANISM: Zea mays
 52 WCPPCR 57
 52 WCPPCR 57
 1 WCGPCK 6
 1 WCGPCK 6
 ISOGAI, TAKAO
SUCIYAMA, TOMOYASU
OTSUKI, TETSUJI
OTSUKI, TETSUJI
WAKAMATSU, AI
SAFO, HIROTUKI
ISHII, SHIZUKO
YAMAMOTO, JUN-ICHI
 4; Conservative
 4; Conservative
 Application US/11195459 No. US20050278803A1
OTSUKA, KAORU
NAGAI, KEIICHI
IRIE, RYOTARO
 Meeley, Robert
Hantke, Sabine
 HIO,
 ISONO, YUUKO
HIO, YURI
 76.6%;
 76.6%;
 1; Mismatches
 Score 36;
Pred. No.
 Score 36;
Pred. No.
 Mismatches
 DB 7;
 Length 128
 Length 128;
 Indels
 Indels
 0
 0;
 Gaps
 Gaps
 0;
 0;
 맑
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 ફ
PRIOR APPLICATION NUMBER: 60/261,014
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j.

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APPLICANT: NAGAHARI, KENJI
APPLICANT: MASCHO, YASUHIKO
TITLE OF INVENTION: NOVEL full length cDNA
FILE REFERENCE: 084335-0191
CURRENT PAPPLICATION NUMBER: US/11/072,512
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: UP 2001-379298
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 4096
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 2145
LENGTH: 129
TYPE: PRT
ORGANISM: Homo sapiens
US-11-072-512-2145
RESULT 55

US-11-054-281-58
; Sequence 58, Application US/11054281
; Publication No. US20060013813A1
; GENERAL INFORMATION:
; APPLICANT: Mezce et al.
; TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same
; FILE REFERENCE: 21402-240CIP
; CURRENT APPLICATION NUMBER: US/11/054,281
; CURRENT FILING DATE: 2005-02-08
; CURRENT FILING DATE: 2005-02-08
 SOFTWARE: PatentIn version 3.3
SEQ ID NO 126
LENGTH: 163
TYPE: PRT
ROANISM: Bacillus licheniformis
US-10-510-386-126
 RESULT 54
US-10-510-386-126
 Matches
 Query Match
Best Local Similarity
 Sequence 126, A
Publication No.
 Matches
 Query Match 76.6%; Score 36; DB 7; Best Local Similarity 100.0%; Pred. No. 15;
 GENERAL INFORMATION:
 APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
 APPLICANT: Andersen, Jens Tonne
APPLICANT: Clausen, Ib Groth
APPLICANT: Jorgensen, Steen Tr
 TITLE OF INVENTION: Improved Bacillus Host Cell
FILE REFERENCE: 10294.204-US
CURRENT PEPLICATION NUMBER: US/10/510,386
CURRENT FILING DATE: 2004-10-04
 APPLICANT: Jorgensen, Steen Troels
APPLICANT: Olsen, Peter Bjarke
APPLICANT: Rasmussen, Michael Dolberg
 CURRENT FILING DATE: 2004-10-04
NUMBER OF SEQ ID NOS: 248
 62 WCPPCR 67
 74 CGPCK 78
 1 WCGPCK 6
 2 CGPCK 6
 5; Conservative
 OTSUKA, MOTOYUKI
 SEKI, NAOHIKO
YOSHIKAWA, TSUTOMU
 TAMECHIKA,
 Application US/10510386 o. US20050244922A1
 Conservative
 76.6%; Score 36; DB 6; 66.7%; Pred. No. 18;
 ICHIRO
 0; Mismatches
 Mismatches
 <u>.</u>
 Length 163;
 Length 129
 Indels
 Indels
 0
 0
 Gaps
 Gaps
 0
 0
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FILING DATE: 2001-01-11

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밁
 Query Match
Best Local Similarity
Watches 5; Conserv
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 RESULT 56
US-11-054-281-59
 밁
 Ś
 ; ORGANISM: Mus musculus
US-11-054-281-58
 US-11-054-281-59
 PRIOR APPLICATION NUMBER: 60/261,018
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/318,410
PRIOR FILING DATE: 2001-09-10
PRIOR APPLICATION NUMBER: 60/261,013
PRIOR APPLICATION NUMBER: 60/261,013
PRIOR APPLICATION NUMBER: 60/261,026
 SEQ ID NO 59
 GENERAL INFORMATION:
 Sequence 59, Application US/11054281
Publication No. US20060013813A1
 Matches
 Query Match
Best Local 9
 SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 58
LENGTH: 354
 CURRENT APPLICATION NUMBER: US/11/054,281
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: 60/261,014
PRIOR FILING DATE: 2001-01-11
 APPLICANT: Mezes et al.
TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same
FILE REFERENCE: 21402-240CIP
 PRIOR FILING DATE: 2001-08-17
PRIOR APPLICATION NUMBER: 10/044,564
PRIOR FILING DATE: 2002-01-11
 PRIOR APPLICATION NUMBER: 60/261,029 PRIOR FILING DATE: 2001-01-11
 PRIOR APPLICATION NUMBER: 10/044,564 PRIOR FILING DATE: 2002-01-11
 PRIOR APPLICATION NUMBER: 60/261,013
PRIOR PILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,026
PRIOR FILING DATE: 2001-01-11
 NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn Ver
 PRIOR
 NUMBER OF SEQ ID NOS: 324
 PRIOR APPLICATION NUMBER: 60/313,170 PRIOR FILING DATE: 2001-08-17
 TYPE: PRT ORGANISM: Rattus norvegicus
 TYPE: PRT
 LENGTH:
 APPLICATION NUMBER: 60/261,029
FILING DATE: 2001-01-11
 APPLICATION NUMBER: 60/318,410 FILING DATE: 2001-09-10
 APPLICATION NUMBER: 60/261,018 FILING DATE: 2001-01-11
 APPLICATION NUMBER: 60/313,170
 FILING DATE:
 144 CGPCK 148
163 CGPCK 167
 2 CGPCK 6
 2 CGPCK 6
 Similarity 5; Conserv
 Conservative
 Conservative
 Ver.
 2001-01-11
 76.6%; Score 36;
100.0%; Pred. No.
tive 0; Mismatcl
 100.0%;
 76.6%; Score 36; DB 7; 100.0%; Pred. No. 29;
 0;
 Mismatches
 Mismatches
 DB 7;
 <u>,,</u>
 0; Indels
 Length 373;
 Length 354;
 Indels
 0
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 Gaps
 0
 0
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```
Sequence 52, Application US/11090439

Publication No. US20050266442A1

GENERAL INFORMATION:

APPLICANT: Squillace, Rachel

APPLICANT: Squillace, Rachel

TITLE OF INVENTION: Immortalized Human Tuberous Sclerosis Null

TITLE OF INVENTION: Angiomyolipoma Cell and Method of Use Thereof

PILE REFERENCE: 24318-502

CURRENT APPLICATION NUMBER: US/11/090,439

CURRENT APPLICATION NUMBER: US/56,344

PRIOR APPLICATION NUMBER: 60/556,344

PRIOR APPLICATION NUMBER: 5004-03-25

NUMBER OF SEQ ID NOS: 62

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 52
 CURRENT APPLICATION NUMBER: US/11/090,439
CURRENT FILING DATE: 2005-03-25
PRIOR APPLICATION NUMBER: 60/556,344
PRIOR FILING DATE: 2004-03-25
NUMBER OF SEQ ID NOS: 62
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 54
LENGTH: 380
TYPE: PRT
ORGANISM: Homo mapiens
 US-11-090-439-54
Sequence 54, Application US/11090439
; Publication No. US20050266442A1
; GENERAL INFORMATION:
APPLICANT: Squillace, Rachel
; APPLICANT: Weiner, Michael P.
; TITLE OF INVENTION: Amgiomyolipoma Cell and Method of Use Thereof
FILE REFERENCE: 24318-502
FILE REFERENCE: 24318-502
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 RESULT 57
US-11-090-439-52
 US-11-054-281-60
 RESULT 59
 US-11-090-439-54
 RESULT 58
 US-11-090-439-52
Sequence 60, Application US/11054281

Publication No. US20060013813A1
GENERAL INFORMATION:
APPLICANT: Mezes et al.
TITLE OF INVENTION: Proteins and Nucleic Acids Encoding
FILE REFERENCE: 21402-240CIP
CURRENT APPLICATION NUMBER: US/11/054,281
 Query Match
Best Local Similarity
 Query Match
 Matches
 Best Local Similarity Matches 5; Conserv
 TYPE: PRT ORGANISM: Homo sapiens
 LENGTH: 380
 170 CGPCK 174
 170 CGPCK 174
 2 CGPCK 6
 2 CGPCK 6
 5
 Conservative
 Conservative
 76.6%; but
/ 100.0%; Pre
 76.6%; Score 36; DB 7; 100.0%; Pred. No. 31;
 0,
 Score 36;
Pred. No.
 Mismatches
 Mismatches
 DB 7;
31;
 0,
 0
 Length 380,
 Length 380;
 Indels
 Same
 <u>.</u>.
 0,
 Gaps
 Gaps
 0
 0;
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CURRENT FILING DATE:

2005-02-08

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RESULT 60
US-10-763-712A-10
; Sequence 10, Application US/10763712A
; Publication No. US20050266541A1
; GENERAL INFORMATION:
; APPLICANT: Solazyme, Inc.
; APPLICANT: Dillon, Harrison F.
RESULT 61
US-10-763-712A-11
; Sequence 11, Application US/10763712A
; Publication No. US20050266541A1
 밁
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 Query Match
Best Local Similarity
Marches 5; Conserva
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 SOFTWARE: PatentIn Ver. 2
; SEQ ID NO 60
; LENGTH: 380
; TYPE: PRT
; ORGANISM: Homo sapiens
 US-10-763-712A-10
 US-11-054-281-60
 PRIOR APPLICATION NUMBER: 60/261,018
PRIOR FILING DATE: 2001-01-11
PRIOR PELICATION NUMBER: 60/318,410
PRIOR PELICATION NUMBER: 60/261,013
PRIOR PELICATION NUMBER: 60/261,013
PRIOR PELICATION NUMBER: 60/261,026
PRIOR PELICATION NUMBER: 60/261,026
PRIOR APPLICATION NUMBER: 60/261,026
PRIOR PILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/261,029
PRIOR FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/313,170
PRIOR PILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 60/313,170
PRIOR PILING DATE: 2002-01-11
PRIOR APPLICATION NUMBER: 10/044,564
PRIOR FILING DATE: 2002-01-11
NUMBER OF SEQ ID NOS: 324
 NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version 3.2
SEQ ID NO 10
 APPLICANT: Dillon, Harrison F.

TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen
TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen
FILE REFERENCE: H2042101-CIP
CURRENT APPLICATION NUMBER: US/10/763,712A
CURRENT FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR FILING DATE: 2002-11-04
PRIOR FILING DATE: 2002-01-04-12
PRIOR APPLICATION NUMBER: US 10/411,910
PRIOR FILING DATE: 2003-09-03
PRIOR APPLICATION NUMBER: US 60/500,032
PRIOR PRIOR DATE: 2003-09-03
PRIOR FILING DATE: 2003-09-03
PRIOR FILING DATE: 2003-09-03
 Matches
 Query Match
 PRIOR APPLICATION NUMBER: 60/261,014 PRIOR FILING DATE: 2001-01-11
 TYPE: PRT ORGANISM: Clostridium perfringens
 LENGTH: 572
 Local Similarity hes 5; Conserv
 170 CGPCK 174
 86
 2 CGPCK 6
 2 CGPCK 6
 CGPCK 102
 Conservative
 Conservative
 76.6%; Score 36; DB 7; 100.0%; Pred. No. 31;
 76.6%; Score 36; DB 6; Length 572; 100.0%; Pred. No. 40;
 0; Mismatches
 0; Mismatches
 0
 0
 Length 380
 Indels
 Indels
 ç
 0,
 Gaps
 0;
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RESULT 63 US-10-763-712A-109

Sequence 109, Application US/10763712A Publication No. US20050266541A1 GENERAL INFORMATION:
APPLICANT: Solazyme, Inc.
APPLICANT: Dillon, Harrison F.

Solazyme, Inc. Dillon, Harrison F.

```
Sequence 37, Application US/10763712A

; Publication No. US20050266541A1

; GENERAL INFORMATION:
 APPLICANT: Solazyme, Inc.

; APPLICANT: Solazyme, Inc.

; APPLICANT: Dillon, Harrison F.

; TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen

; TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen

; TITLE REFERINCE: H2042101-CIP

; CURRENT APPLICATION NUMBER: US/10/763,712A

; CURRENT FILING DATE: 2004-01-21

; PRIOR APPLICATION NUMBER: US 10/287,750

; PRIOR APPLICATION NUMBER: US 10/287,750

; PRIOR FILING DATE: 2003-04-12

; PRIOR APPLICATION NUMBER: US 00/500,032

; PRIOR APPLICATION NUMBER: US 60/500,032

; PRIOR FILING DATE: 2003-09-03

; NUMBER OF SEQ ID NOS: 184
 APPLICANT: Solazyme, Inc.
APPLICANT: Dillon, Harrison F.
APPLICANT: Dillon, Harrison F.
TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen
TITLE OF INVENTION: Production
FILE REFERENCE: H2042101-CIP
CURRENT APPLICATION NUMBER: US/10/763,712A
CURRENT FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR FILING DATE: 2002-11-04
PRIOR FILING DATE: 2003-04-12
PRIOR APPLICATION NUMBER: US 60/500,032
PRIOR FILING DATE: 2003-04-12
PRIOR FILING DATE: 2003-09-03
NUMBER OF SEQ ID NOS: 184
SOFTWARE: Patentin version 3.2
SEQ ID NO 1:
TENCTH: 572
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 RESULT 62
US-10-763-712A-37
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 US-10-763-712A-37
 US-10-763-712A-11
 SOFTWARE: PatentIn version 3.2 SEQ ID NO 37
 Matches
 Query Match
Best Local Similarity
 Matches
 Query Match
 GENERAL INFORMATION:
 Query Match 76.6%; Score 36; DB Best Local Similarity 100.0%; Pred. No. 40;
 LENGTH: 572
TYPE: PRT
ORGANISM: Clostridium perfringens
 TYPE: PRT Clostridium perfringens
 LENGTH: 572
 98 CGPCK 102
98 CGPCK 102
 2 CGPCK 6
 2 CGPCK 6
 5; Conservative
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 Conservative
 100.0%;
 76.6%; Score 36; DB 6; 100.0%; Pred. No. 40;
 0; Mismatches
 0; Mismatches
 6
 0;
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 Length 572;
 Length 572;
 Indels
 0
 0
 Gaps
 Gaps
 0;
 0;
```

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PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR FILING DATE: 2002-11-04
PRIOR PELICATION NUMBER: US 10/411,910
PRIOR PELICATION NUMBER: US 60/500,032
PRIOR APPLICATION NUMBER: US 60/500,032
PRIOR FILING DATE: 2003-09-03
NUMBER OF SEQ ID NOS: 184
SOFTWARE: PATENTIN VERBION 3.2
SEQ ID NO 109
SEQ ID NO 109
LENGTH: 572
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 ; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 110
; LENGTH: 572
; TYPE: PRT
; ORGANISM: C. perfringens
US-10-763-712A-110
 APPLICANT: Solazyme, Inc.
APPLICANT: Solazyme, Inc.
APPLICANT: Dillon, Harrison F.
APPLICANT: Dillon, Harrison F.
TITLE OF INVENTION: Methods and Compositions
TITLE OF INVENTION: Production
FILE REFERENCE: H2042101-CIP
CURRENT APPLICATION NUMBER: US/10/763,712A
CURRENT FILING DATE: 2004-01-21
PRIOR APPLICATION NUMBER: US 10/287,750
PRIOR FILING DATE: 2002-11-04
PRIOR FILING DATE: 2002-10-04
PRIOR FILING DATE: 2003-09-03
PRIOR FILING DATE: 2003-09-03
PRIOR PILING DATE: 2003-09-03
PRIOR PILING DATE: 2003-09-03
PRIOR PILING DATE: 2003-09-03
NUMBER OF SEQ ID NOS: 184
 밁
 US-10-055-877-255
 RESULT 65
 밁
 S
 US-10-763-712A-110
 US-10-763-712A-109
 Sequence 255, Application US/1005877
Publication No. US20050288241A1
GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
APPLICANT: Tcherney, Velizar
 Query Match
Best Local
 Query Match
Best Local (
 GENERAL INFORMATION:
 Sequence 110, Application US/10763712A Publication No. US20050266541A1
 Matches
 Matches
 CURRENT APPLICATION NUMBER: US/10/763,712A CURRENT FILING DATE: 2004-01-21
 TITLE OF INVENTION: Methods and Compositions for Evolving Microbial Hydrogen
 FILE REFERENCE: H2042101-CIP
 APPLICANT:
 ORGANISM: C. perfringens
 98 CGPCK 102
 98 CGPCK 102
 l Similarity
5; Conserv
 2 CGPCK 6
 2 CGPCK 6
 h 76.6%; Score 36; DB Similarity 100.0%; Pred. No. 40; S; Conservative 0; Mismatches
 Conservative
 100.0%;
 76.6%; Score 36; 100.0%; Pred. No.
 0;
 Compositions for Evolving Microbial Hydrogen
 Mismatches
 DB 6;
 DB 6;
 <u>.</u>
 0
 Length 572;
 Length 572;
 Indels
 Indels
 <u>,</u>
 0;
 0
 0;
 ; TYPE: PRT; ORGANISM: Homo sapiens US-10-055-877-255
 RESULT 66
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 5
 US-10-055-877-254
 CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION UNMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR FILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR APPLICATION NUMBER: 60/264,139
Sequence 254, Application US/1005877
Publication No. US20050288241A1
GENERAL INFORMATION:
APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
 Query Match
Best Local Similarity
Matches 5; Conserv
 SEQ ID NO 255
LENGTH: 654
 APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
 APPLICANT:
APPLICANT:
APPLICANT:
 Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 512 SOFTWARE: PatentIn Ver. 2.1
 PRIOR FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR FILING DATE: 2001-03-14
 APPLICANT:
 APPLICANT:
 PRIOR APPLICATION NUMBER: 60/275,927 PRIOR FILING DATE: 2001-03-14
 PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 60/272,870
 PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,478
 APPLICANT: Boldog, Ference
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
 PRIOR APPLICATION NUMBER: 60/263,351
 PRIOR FILING DATE: 2001-01-26
 APPLICANT:
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 247 CGPCK 251
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Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
 Kekuda, Ramesh
Guo, Xiaojia
Zerhusen, Bryan
Andrew, David
 Andrew, David
Mezes, Peter
 Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
 Shenoy, Suresh
 Wolenc, Adam
Baumgartner, Jason
Shimkets, Richard
 Patturajan, Meera
Burgess, Cahterine
 Anderson, David
 Eisen, Andrew
 Ratelli, Luca
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Gaps

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Miller, Charles

Velizar

Tchernev, Zhong, Mei

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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-254
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Sequence 4, Application US/10831997
Publication No. US20050244834A1
 SOFTWARE: PatentIn Ver. SEQ ID NO 254
 Remaining Prior Application data removed - See File Wrapper or PALM.
 CURRENT APPLICATION NUCLEUR FILING DATE:
 PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
 RIOR FILING DATE:
 RIOR APPLICATION NUMBER: 60/263,598
 PHICANT: Boldog, Ference
TILE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
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mes 5; Conserv
 FILING DATE: 2001-01-24
APPLICATION NUMBER: 60/264,117
FILING DATE: 2001-01-25
 APPLICATION NUMBER: 60/263,799
FILING DATE: 2001-01-24
 APPLICATION NUMBER: 60/275,927
 FILING DATE: 2001-03-14
 APPLICATION NUMBER: 60/272,870 FILING DATE: 2001-03-02
 FILING DATE
 APPLICATION NUMBER: 60/264,139
 APPLICATION NUMBER: 60/275,990
 APPLICATION NUMBER: 60/263,351
 APPLICATION NUMBER: 60/264,478
 249 CGPCK 253
 APPLICATION NUMBER: US/10/055,877
 LING DATE: 2001-01-25
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Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
 Guo, Xiaojia
Zerhusen, Bryan
Andrew, David
 Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
 Shenoy, Suresh
 Gusev, Vladimir
 Mezes, Peter
Patturajan, Meera
Burgess, Cahterine
 Casman, Stacie
 Baumgartner, Jason
Shimkets, Richard
 Burgess, Cahte
Eisen, Andrew
 Ratelli, Luca
Kekuda, Ramesh
 Volenc, Adam
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 2001-01-23
 2001-01-30
 2001-01-26
 2001-01-25
 2002-01-22
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Pred. No. 44;
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APPLICANT: Daley, George Q.

APPLICANT: McCarthy, Jeanette J.

TITLE OF INVENTION: SINGLE NUCLECTIDE POLYMORPHISMS IN GENES FILE REFERENCE: 2825.1027-001

CURRENT APPLICATION NUMBER: US/10/831,997

CURRENT FILLING DATE: 2004-04-26

PRIOR APPLICATION NUMBER: US/99/657,472

PRIOR FILLING DATE: 2000-09-07

PRIOR FILLING DATE: 2000-09-07

PRIOR APPLICATION NUMBER: US 60/153,357

PRIOR APPLICATION NUMBER: US 60/153,357

PRIOR APPLICATION NUMBER: US 60/25,724

PRIOR APPLICATION NUMBER: US 60/225,724

PRIOR FILING DATE: 2000-08-16

PRIOR FILING DATE: 2000-08-26

PRIOR FILING DATE: 2000-08-26
 RESULT 68
US-10-055-877-252
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SEQ ID NO 4
LENGTH: 961
TYPE: PRT
ORGANISM: Homo sapiens
US-10-831-997-4
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TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
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Pena, Carol
 Baumgartner, Jason
Shimkets, Richard
Gusev, Vladimir
 Patturajan, Meera
Burgess, Cahterine
 Anderson, David
Ballinger, Rober
 Padigaru, Muralidhara
Miller, Charles
 Casman, Stacie
 Shenoy, Suresh
 Eisen, Andrew
Wolenc, Adam
 Ratelli, Luca
Kekuda, Ramesh
 Spytek, Kimberly
 Gerlach,
 Zhong, Mei
 Chernev,
 Valerie
 Velizar
 DB 6;
 Length 961;
 Indels
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 Gaps
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US-10-055-877-252
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 Sequence 73, Application US/10055877 Publication No. US20050288241A1
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Best Local (
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 Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 512
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 APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
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 APPLICATION NUMBER: 60/263,799
FILING DATE: 2001-01-23
ADDITE: 2001-01-24
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 FILING DATE:
 APPLICATION NUMBER: 60/272,870
 NT FILING DATE: 2002-01-22
APPLICATION NUMBER: 60/262,892
 FILING DATE: 2001-01-25
 APPLICATION NUMBER: 60/264,117
 APPLICATION NUMBER: 60/263,598
 FILING DATE: 2001-03-02
APPLICATION NUMBER: 60/275,990
 FILING DATE:
 APPLICATION NUMBER: 60/264,139
 APPLICATION NUMBER: 60/275,927
 FILING DATE:
 APPLICATION NUMBER: 60/263,351
 FILING DATE:
 APPLICATION NUMBER: 60/264,478
 FILING DATE:
 65 CGPCK 69
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Vernet, Corine
Taupier Jr., Raymond
Pena, Carol
 Ballinger, Robert
Gerlach, Valerie
Spytek, Kimberly
 Padigaru, muraii
Miller, Charles
Boldog, Ference
 Casman, Stacie
 Shenoy, Suresh
 Wolenc, Adam
Baumgartner, Jason
Shimkets, Richard
 Eisen, Andrew
 Patturajan, Meera
Burgess, Cahterine
 Mezes, Peter
 Zerhusen, Brya
Andrew, David
 Guo, Xiaojia
Zerhusen, Bryan
 Ratelli, Luca
Kekuda, Ramesh
 Anderson, David
 Tchernev,
 Conservative
 Mei
 2001-03-14
 2001-01-26
 2001-01-25
 2001-03-14
 2001-01-30
 2001-01-19
 76.6%;
100.0%;
 Velizar
 Robert
 0
 Score 36;
Pred. No.
 Mismatches
 DB 6;
 Length 1857;
 Indels
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 US-10-055-877-253
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PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR FILING DATE: 2001-01-25
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Matches 5
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 Sequence 253, P
Publication No.
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 FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
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 APPLICANT: DeCristofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
 Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 512
 PRIOR FILING DATE: 2001-03-14
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PRIOR FILING DATE: 2001-03-14
 PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-30
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 LENGTH: 2084
 FILING DATE: 2001-03-02
APPLICATION NUMBER: 60/275,990
 APPLICATION NUMBER: 60/272,870
 APPLICATION NUMBER: 60/264,478
 317 CGPCK 321
 2 CGPCK 6
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 Wolenc, Adam
Baumgartner, Jason
Shimkets, Richard
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Vernet, Corine
Taupier Jr., Raymond
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 Zerhusen, Zavid
 Gerlach, Valerie
Spytek, Kimberly
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 Guo, Xiaojia
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Kekuda, Ramesh
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 Zhong, Mei
 Tchernev,
 Pena, Carol
 Application US/10055877
b. US20050288241A1
 Conservative
 Velizar
 76.6%; Score 36;
100.0%; Pred. No.
tive 0; Mismatch
 Mismatches
 DB 6;
92;
 Length 2084;
 Indels
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 Gaps
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APPLICANT: Li, Li
APPLICANT: Boldog, Ference
APPLICANT: Boldog, Ference
PRICANT: Boldog, Ference
FILE REFERENCE: 21402-221
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT PILING DATE: 2002-01-22
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FRIOR APPLICATION NUMBER: 60/263,598
FRIOR APPLICATION NUMBER: 60/263,598
FRIOR APPLICATION NUMBER: 60/263,799
FRIOR FILING DATE: 2001-01-24
FRIOR FILING DATE: 2001-01-25
FRIOR APPLICATION NUMBER: 60/264,478
FRIOR FILING DATE: 2001-01-31-4
FRIOR APPLICATION NUMBER: 60/275,990
FRIOR FILING DATE: 2001-03-14
FRIOR
 Query Match 76.6%; Score 36; DB 6; Length 2098; Best Local Similarity 100.0%; Pred. No. 93; Matches 5; Conservative 0; Mismatches 0; Indels
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Search completed: February 23, 2006, 00:48:12 Job time : 15.5 secs

267 CGPCK 271

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